

# Schema documentation for no-specific-site.xsd

6 september 2010

## Table of Contents

Namespace: "http://purl.org/obbr/pathology/cancer/" .....	6
Schemas .....	6
Main schema no-specific-site.xsd .....	6
Included schema valuesets.xsd .....	6
Included schema values.xsd .....	6
Included schema generic.xsd .....	6
Elements .....	6
Element diagnosis .....	6
Element obbr:generic.content / obbr:specify .....	8
Element obbr:generic.content / obbr:label .....	8
Element obbr:generic.content / obbr:comment .....	8
Element history.section / history .....	9
Element history.clinicalFinding.item / clinicalFinding .....	10
Element history.clinicalFinding.response / response .....	11
Element history.pathologicFinding.item / pathologicFinding .....	11
Element history.pathologicFinding.response / response .....	12
Element history.priorTherapy.item / priorTherapy .....	13
Element history.priorTherapy.response / response .....	14
Element history.radiologicFinding.item / radiologicFinding .....	15
Element history.radiologicFinding.response / response .....	16
Element specimen.section / specimen .....	17
Element specimen.procedure.item / procedure .....	19
Element specimen.procedure.response / response .....	19
Element specimen.size.item / size .....	20
Element obbr:size / obbr:response .....	21
Element specimen.weight.item / weight .....	22
Element specimen.weight.response / response .....	23
Element specimen.site.item / site .....	23
Element specimen.site.response / response .....	24
Element specimen.integrity.item / integrity .....	25
Element specimen.integrity.response / response .....	26
Element tumor.section / tumor .....	27
Element tumor.focus.section / focus .....	28
Element tumor.site.item / site .....	30
Element tumor.site.response / response .....	30
Element tumor.size.item / size .....	31
Element tumor.histologicType.item / histologicType .....	32
Element tumor.histologicType.response / response .....	33
Element tumor.grade.item / grade .....	34
Element tumor.grade.response / response .....	35
Element extent.section / extent .....	36
Element extent.contiguousExtension.section / contiguousExtension .....	37
Element extent.contiguousExtension.status.section / status .....	38
Element extent.contiguousExtension.status.result.item / result .....	39
Element extent.contiguousExtension.status.result.response / response .....	40
Element extent.contiguousExtension.status.distance.item / distance .....	41
Element obbr:length / obbr:response .....	42
Element accessory.section / accessory .....	43
Element accessory.treatmentResponse.item / treatmentResponse .....	44
Element accessory.treatmentResponse.response / response .....	45
Element accessory.lymphvascularInvasion.item / lymphvascularInvasion .....	46
Element accessory.lymphvascularInvasion.response / response .....	47
Element accessory.perineuralInvasion.item / perineuralInvasion .....	48
Element accessory.perineuralInvasion.response / response .....	49
Element margins.section / margins .....	50
Element margins.margin.section / margin .....	51
Element margins.margin.status.section / status .....	52
Element margins.margin.status.result.item / result .....	54
Element margins.margin.status.result.response / response .....	54
Element margins.margin.distance.item / distance .....	55

Element nodes.section / nodes .....	56
Element nodes.status.section / status .....	57
Element nodes.status.result.section / result .....	59
Element nodes.status.result.response / response .....	60
Element nodes.status.largestMetastasis.item / largestMetastasis .....	60
Element nodes.status.extracapsularExtension.item / extracapsularExtension .....	61
Element nodes.extracapsularExtension.response / response .....	62
Element distant.section / distant .....	63
Element distant.status.section / status .....	64
Element distant.status.result.item / result .....	66
Element distant.status.result.response / response .....	67
Element studies.section / studies .....	67
Element studies.section.content / study .....	68
Element studies.study.item / study .....	69
Element studies.study.response / response .....	70
Element stage.section / stage .....	71
Element stage.prefix.item / prefix .....	73
Element stage.prefix.response / response .....	74
Element stage.multiplicity.item / multiplicity .....	74
Element stage.multiplicity.response / response .....	75
Element stage.pT.item / pT .....	76
Element stage.pT.response / response .....	76
Element stage.pN.item / pN .....	77
Element stage.pN.response / response .....	78
Element stage.pM.item / pM .....	79
Element stage.pM.response / response .....	80
Element stage.group.item / group .....	81
Element stage.group.response / response .....	82
Element additional.section / additional .....	83
Element additional.finding.item / finding .....	84
Element additional.finding.response / response .....	85
Simple Types .....	86
Simple Type obbr:history.clinicalFinding.valueset .....	86
Simple Type obbr:date.valueset .....	86
Simple Type obbr:history.pathologicFinding.valueset .....	87
Simple Type obbr:history.priorTherapy.valueset .....	87
Simple Type obbr:history.radiologicFinding.valueset .....	87
Simple Type obbr:laterality.valueset .....	87
Simple Type obbr:specimen.procedure.response.valueset .....	88
Simple Type obbr:scalar.valueset .....	88
Simple Type obbr:lengthUnit.valueset .....	88
Simple Type obbr:relation.valueset .....	88
Simple Type obbr:weightUnit.valueset .....	89
Simple Type obbr:specimen.site.valueset .....	89
Simple Type obbr:specimen.integrity.valueset .....	89
Simple Type obbr:tumor.site.valueset .....	89
Simple Type obbr:tumor.grade.valueset .....	89
Simple Type obbr:tumor.gradeSystem.valueset .....	90
Simple Type obbr:tumor.focus.type.valueset .....	90
Simple Type obbr:focality.valueset .....	90
Simple Type obbr:qualitative.valueset .....	90
Simple Type obbr:extent.contiguousExtension.status.result.for.valueset .....	91
Simple Type obbr:extent.contiguousExtension.status.distance.for.valueset .....	91
Simple Type obbr:base.values .....	91
Simple Type obbr:extent.status.of.valueset .....	91
Simple Type obbr:accessory.treatmentResponse.valueset .....	92
Simple Type obbr:accessory.lymphvascularInvasion.valueset .....	92
Simple Type obbr:accessory.perineuralInvasion.valueset .....	92
Simple Type obbr:margins.margin.status.result.for.valueset .....	92
Simple Type obbr:margins.margin.distance.for.valueset .....	93
Simple Type obbr:margins.margin.status.of.valueset .....	93
Simple Type obbr:tally.valueset .....	93
Simple Type obbr:nodes.status.result.for.valueset .....	93
Simple Type obbr:nodes.status.result.type.valueset .....	93
Simple Type obbr:nodes.status.of.valueset .....	94
Simple Type obbr:nodes.status.technique.valueset .....	94
Simple Type obbr:distant.status.result.for.valueset .....	94
Simple Type obbr:distant.status.of.valueset .....	94
Simple Type obbr:stage.prefix.valueset .....	95
Simple Type obbr:multiplicity.valueset .....	95
Simple Type obbr:stage.pT.valueset .....	95
Simple Type obbr:stage.pN.valueset .....	95

Simple Type obbr:stage.pM.valueset .....	95
Simple Type obbr:stage.group.valueset .....	96
Simple Type obbr:additional.finding.valueset .....	96
Simple Type obbr:additional.finding.of.valueset .....	96
Simple Type obbr:margins.margin.status.focality.valueset .....	96
Simple Type obbr:margins.margin.distance.from.valueset .....	96
Simple Type obbr:nodes.status.extracapsular.valueset .....	97
Simple Type obbr:tumor.histologicType.valueset .....	97
Simple Type obbr:empty .....	97
Simple Type obbr:accessory.treatmentResponse.values .....	97
Simple Type obbr:extent.contiguousExtension.status.result.for.values .....	98
Simple Type obbr:extent.contiguousExtension.status.distance.for.values .....	98
Simple Type obbr:history.priorTherapy.values .....	98
Simple Type obbr:history.pathologicFinding.values .....	98
Simple Type obbr:history.clinicalFinding.values .....	98
Simple Type obbr:history.radiologicFinding.values .....	99
Simple Type obbr:margins.margin.status.result.for.values .....	99
Simple Type obbr:distant.status.of.values .....	99
Simple Type obbr:distant.status.result.for.values .....	99
Simple Type obbr:nodes.status.result.for.values .....	100
Simple Type obbr:nodes.status.result.type.values .....	100
Simple Type obbr:nodes.status.technique.values .....	100
Simple Type obbr:specimen.integrity.values .....	101
Simple Type obbr:stage.prefix.values .....	101
Simple Type obbr:stage.pT.values .....	101
Simple Type obbr:stage.pN.values .....	101
Simple Type obbr:stage.pM.values .....	102
Simple Type obbr:stage.group.values .....	102
Simple Type obbr:tumor.grade.values .....	102
Simple Type obbr:tumor.gradeSystem.values .....	102
Simple Type obbr:tumor.focus.type.values .....	103
Simple Type obbr:tumor.site.values .....	103
Simple Type obbr:positiveDecimal .....	103
Simple Type obbr:focality.values .....	103
Simple Type obbr:laterality.values .....	104
Simple Type obbr:lengthUnit.values .....	104
Simple Type obbr:multiplicity.values .....	104
Simple Type obbr:qualitative.values .....	105
Simple Type obbr:relation.values .....	105
Simple Type obbr:weightUnit.values .....	105
Element Group .....	106
Element Group obbr:generic.content .....	106
Element Group diagnosis.section.content .....	106
Element Group history.section .....	108
Element Group history.section.content .....	108
Element Group history.clinicalfinding.item .....	109
Element Group history.clinicalFinding.response .....	109
Element Group history.pathologicFinding.item .....	110
Element Group history.pathologicFinding.response .....	110
Element Group history.priorTherapy.item .....	111
Element Group history.priorTherapy.response .....	111
Element Group history.radiologicFinding.item .....	111
Element Group history.radiologicFinding.response .....	112
Element Group specimen.section .....	112
Element Group specimen.section.content .....	112
Element Group specimen.procedure.item .....	113
Element Group specimen.procedure.response .....	114
Element Group specimen.size.item .....	114
Element Group specimen.size.response .....	114
Element Group obbr:size .....	115
Element Group specimen.weight.item .....	115
Element Group specimen.weight.response .....	115
Element Group specimen.site.item .....	116
Element Group specimen.site.response .....	116
Element Group specimen.integrity.item .....	116
Element Group specimen.integrity.response .....	117
Element Group tumor.section .....	117
Element Group tumor.section.content .....	117
Element Group tumor.focus.section .....	118
Element Group tumor.site.item .....	118
Element Group tumor.site.response .....	119
Element Group tumor.size.item .....	119

Element Group tumor.size.response .....	119
Element Group tumor.histologicType.item .....	120
Element Group tumor.histologicType.response .....	120
Element Group tumor.grade.item .....	120
Element Group tumor.grade.response .....	121
Element Group extent.section .....	121
Element Group extent.section.content .....	121
Element Group extent.contiguousExtension.section .....	122
Element Group extent.contiguousExtension.status.section .....	122
Element Group extent.contiguousExtension.status.result.item .....	123
Element Group extent.contiguousExtension.status.result.response .....	123
Element Group extent.contiguousExtension.status.distance.item .....	123
Element Group obbr:length .....	124
Element Group accessory.section .....	124
Element Group accessory.section.content .....	124
Element Group accessory.treatmentResponse.item .....	125
Element Group accessory.treatmentResponse.response .....	125
Element Group accessory.lymphvascularInvasion.item .....	126
Element Group accessory.lymphvascularInvasion.response .....	126
Element Group accessory.perineuralInvasion.item .....	126
Element Group accessory.perineuralInvasion.response .....	127
Element Group margins.section .....	127
Element Group margins.section.content .....	128
Element Group margins.margin.section .....	128
Element Group margins.margin.section.content .....	128
Element Group margins.margin.status.section .....	129
Element Group margins.margin.status.result.item .....	129
Element Group margins.margin.status.result.response .....	129
Element Group margins.margin.distance.item .....	130
Element Group nodes.section .....	130
Element Group nodes.section.content .....	130
Element Group nodes.status.section .....	131
Element Group nodes.status.result.section .....	131
Element Group nodes.status.result.response .....	132
Element Group nodes.status.largestMetastasis.item .....	132
Element Group nodes.status.extracapsularExtension.item .....	132
Element Group nodes.extracapsularExtension.response .....	133
Element Group distant.section .....	133
Element Group distant.section.content .....	133
Element Group distant.status.section .....	134
Element Group distant.status.section.content .....	134
Element Group distant.status.result.item .....	134
Element Group distant.status.result.response .....	135
Element Group studies.section .....	135
Element Group studies.section.content .....	135
Element Group studies.study.item .....	136
Element Group studies.study.response .....	136
Element Group stage.section .....	137
Element Group stage.section.content .....	137
Element Group stage.prefix.item .....	138
Element Group stage.prefix.response .....	138
Element Group stage.multiplicity.item .....	138
Element Group stage.multiplicity.response .....	139
Element Group stage.pT.item .....	139
Element Group stage.pT.response .....	139
Element Group stage.pN.item .....	140
Element Group stage.pN.response .....	140
Element Group stage.pM.item .....	141
Element Group stage.pM.response .....	141
Element Group stage.group.item .....	141
Element Group stage.group.response .....	142
Element Group additional.section .....	142
Element Group additional.section.content .....	142
Element Group additional.finding.item .....	143
Element Group additional.finding.response .....	143
Attribute Groups .....	143
Attribute Group obbr:generic.attrib .....	143
Attribute Group obbr:date .....	144
Attribute Group obbr:quantity .....	145
Attribute Group obbr:weight .....	145
Attribute Group obbr:tally .....	145
Namespace: "http://www.w3.org/XML/1998/namespace" .....	146

Schemas .....	146
Imported schema xmlid.xsd .....	146
Attributes .....	146
Attribute @xml:id .....	146
Namespace: "" .....	146
Attributes .....	146
Attribute obbr:generic.content / @value .....	146
Attribute history.clinicalFinding.response / response / @value .....	146
Attribute obbr:date / @date .....	147
Attribute obbr:date / @start .....	147
Attribute obbr:date / @end .....	147
Attribute obbr:date / @duration .....	147
Attribute history.pathologicFinding.response / response / @value .....	147
Attribute history.priorTherapy.response / response / @value .....	147
Attribute history.radiologicFinding.response / response / @value .....	148
Attribute obbr:generic.content / @on .....	148
Attribute specimen.procedure.response / response / @on .....	148
Attribute specimen.procedure.response / response / @value .....	148
Attribute obbr:quantity / @value .....	148
Attribute obbr:quantity / @unit .....	148
Attribute obbr:quantity / @relation .....	148
Attribute obbr:weight / @value .....	149
Attribute obbr:weight / @unit .....	149
Attribute obbr:weight / @relation .....	149
Attribute specimen.site.response / response / @value .....	149
Attribute specimen.integrity.response / response / @value .....	149
Attribute tumor.site.response / response / @on .....	149
Attribute tumor.site.response / response / @value .....	150
Attribute tumor.histologicType.response / response / @value .....	150
Attribute tumor.grade.response / response / @value .....	150
Attribute tumor.grade.item / grade / @system .....	150
Attribute tumor.focus.section / focus / @type .....	150
Attribute obbr:generic.content / @extent .....	150
Attribute extent.contiguousExtension.status.result.response / response / @extent .....	151
Attribute extent.contiguousExtension.status.result.response / response / @value .....	151
Attribute extent.contiguousExtension.status.result.item / result / @for .....	151
Attribute extent.contiguousExtension.status.result.item / result / @on .....	151
Attribute extent.contiguousExtension.status.distance.item / distance / @for .....	151
Attribute extent.contiguousExtension.status.distance.item / distance / @from .....	151
Attribute extent.contiguousExtension.status.section / status / @of .....	152
Attribute accessory.treatmentResponse.response / response / @value .....	152
Attribute accessory.lymphvascularInvasion.response / response / @value .....	152
Attribute accessory.perineuralInvasion.response / response / @value .....	152
Attribute obbr:generic.content / @closest .....	152
Attribute margins.margin.status.result.response / response / @closest .....	152
Attribute margins.margin.status.result.response / response / @extent .....	153
Attribute margins.margin.status.result.response / response / @value .....	153
Attribute margins.margin.status.result.item / result / @for .....	153
Attribute margins.margin.distance.item / distance / @for .....	153
Attribute margins.margin.status.section / status / @of .....	153
Attribute margins.margin.status.section / status / @on .....	153
Attribute obbr:tally / @value .....	154
Attribute obbr:tally / @relation .....	154
Attribute nodes.status.result.section / result / @for .....	154
Attribute nodes.status.result.section / result / @type .....	154
Attribute nodes.extracapsularExtension.response / response / @value .....	154
Attribute nodes.status.section / status / @of .....	154
Attribute nodes.status.section / status / @on .....	155
Attribute nodes.status.section / status / @technique .....	155
Attribute distant.status.response / response / @value .....	155
Attribute distant.status.result.item / result / @for .....	155
Attribute distant.status.section / status / @of .....	155
Attribute distant.status.section / status / @on .....	155
Attribute studies.study.response / response / @value .....	156
Attribute stage.prefix.response / response / @value .....	156
Attribute stage.multiplicity.response / response / @value .....	156
Attribute stage.pT.response / response / @value .....	156
Attribute stage.pN.response / response / @value .....	156
Attribute stage.pM.response / response / @value .....	156
Attribute stage.group.response / response / @value .....	157
Attribute additional.finding.response / response / @value .....	157
Attribute additional.finding.item / finding / @of .....	157

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Attribute diagnosis / @schemaLocation .....	157
Attribute diagnosis / @version .....	157

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

### Schemas

#### Main schema no-specific-site.xsd

Namespace	http://purl.org/obbr/pathology/cancer/
Properties	attribute form default: unqualified element form default: qualified
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

#### Included schema valuesets.xsd

Namespace	http://purl.org/obbr/pathology/cancer/
Properties	attribute form default: unqualified element form default: qualified
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

#### Included schema values.xsd

Namespace	http://purl.org/obbr/pathology/cancer/
Properties	attribute form default: unqualified element form default: qualified
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

#### Included schema generic.xsd

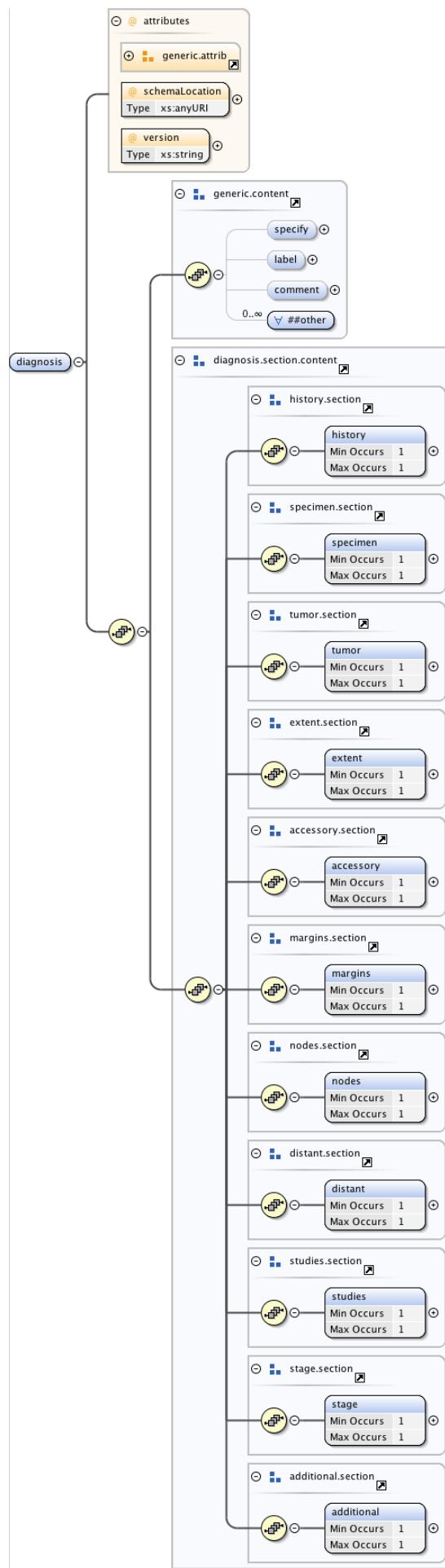
Namespace	http://purl.org/obbr/pathology/cancer/
Properties	attribute form default: unqualified element form default: qualified
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Elements

#### Element diagnosis

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



Properties	content: complex				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , history , specimen , tumor , extent , accessory , margins , nodes , distant , studies , stage , additional				
Children	accessory, additional, distant, extent, history, margins, nodes, obbr:comment, obbr:label, obbr:specify, specimen, stage, studies, tumor				
Instance	<diagnosis xml:id="" schemaLocation="" version=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </diagnosis>				
Attributes	QName	Type	Fixed	Default	Use
	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>				
	<b>schemaLocation</b>	xs:anyURI			required
	<b>version</b>	xs:string			required
	<b>xml:id</b>	xs:ID			optional
Source	<xs:element name="diagnosis"> <xs:complexType> <xs:sequence> <xs:group ref="generic.content"/> <xs:group ref="diagnosis.section.content"/> </xs:sequence> <xs:attributeGroup ref="generic.attribs"/> <xs:attribute name="schemaLocation" type="xs:anyURI" use="required"/> <xs:attribute name="version" type="xs:string" use="required"/> </xs:complexType> </xs:element>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd				

### Element obbr:generic.content / obbr:specify

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	xs:string
Properties	content: simple minOccurs: 0
Source	<xs:element minOccurs="0" name="specify" type="xs:string"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Element obbr:generic.content / obbr:label

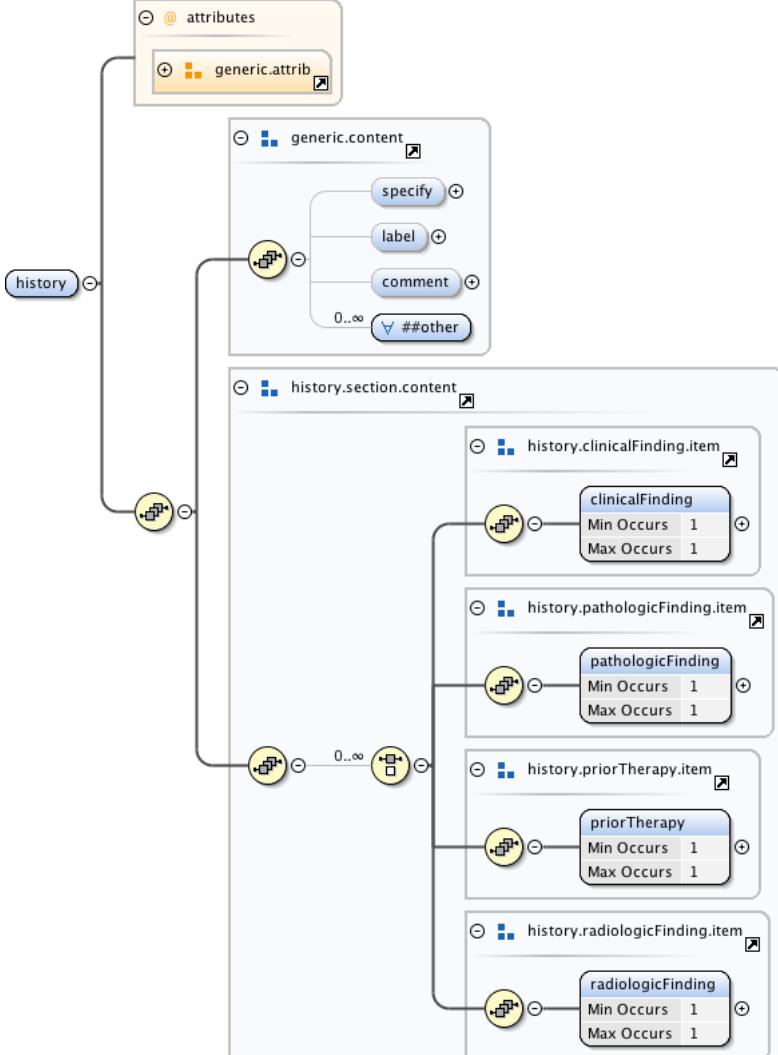
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	xs:string
Properties	content: simple minOccurs: 0
Source	<xs:element minOccurs="0" name="label" type="xs:string"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Element obbr:generic.content / obbr:comment

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Type	xs:string
Properties	<p>content: simple</p> <p>minOccurs: 0</p>
Source	<xs:element minOccurs="0" name="comment" type="xs:string"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

## Element history.section / history

Namespace	http://purl.org/obbr/pathology/cancer/										
Diagram											
Properties	content: complex										
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , ((clinicalFinding)   (pathologicFinding)   (priorTherapy)   (radiologicFinding))										
Children	clinicalFinding, obbr:comment, obbr:label, obbr:specify, pathologicFinding, priorTherapy, radiologicFinding										
Instance	<pre>&lt;history xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/history&gt;</pre>										
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace</td> <td></td> <td></td> <td></td> <td></td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace				
QName	Type	Fixed	Default	Use							
ANY attribute from ANY namespace											

	QName	Type	Fixed	Default	Use
	<b>OTHER than</b> 'http://purl.org/ obbr/pathology/ cancer/'				
	<b>xml:id</b>	xs:ID			optional
Source	<xs:element name="history"> <xs:complexType> <xs:sequence> <xs:group ref="generic.content"/> <xs:group ref="history.section.content"/> </xs:sequence> <xs:attributeGroup ref="generic.attrib"/> </xs:complexType> </xs:element>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd				

**Element history.clinicalFinding.item / clinicalFinding**

Namespace	http://purl.org/obbr/pathology/cancer/																																			
Diagram	<pre> classDiagram     class clinicalFinding     class attributes {         date         genericattrib     }     class genericContent {         specify         label         comment         #other     }     class historyClinicalFindingResponse {         response         Min Occurs 1         Max Occurs 1     }      clinicalFinding "1" -- "*" attributes     clinicalFinding "1" -- "*" genericContent     clinicalFinding "1" -- "*" historyClinicalFindingResponse   </pre>																																			
Properties	content: complex																																			
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																																			
Children	obbr:comment, obbr:label, obbr:specify, response																																			
Instance	<pre> &lt;clinicalFinding date="" duration="" end="" xml:id="" start=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/clinicalFinding&gt;   </pre>																																			
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>date</b></td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>duration</b></td> <td>xs:duration</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>end</b></td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>start</b></td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>					<b>date</b>	obbr:date.valueset			optional	<b>duration</b>	xs:duration			optional	<b>end</b>	obbr:date.valueset			optional	<b>start</b>	obbr:date.valueset			optional	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																																
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<b>end</b>	obbr:date.valueset			optional																																
<b>start</b>	obbr:date.valueset			optional																																
<b>xml:id</b>	xs:ID			optional																																
Source	<xs:element name="clinicalFinding">																																			

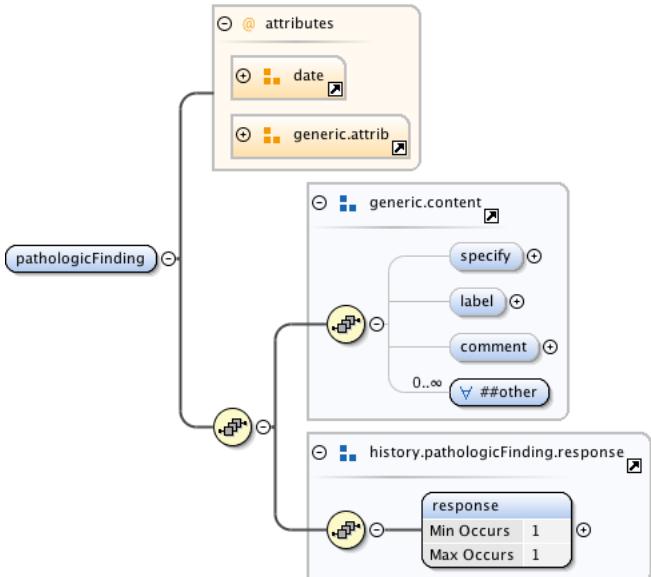
	<pre> &lt;xs:complexType&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:group ref="history.clinicalFinding.response"/&gt;   &lt;/xs:sequence&gt;   &lt;xs:attributeGroup ref="date"/&gt;   &lt;xs:attributeGroup ref="generic.attrib"/&gt; &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element history.clinicalFinding.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/																				
Diagram	<pre> graph TD     response[response] --&gt; attributes[attributes]     attributes --&gt; value["@ value Type: history.clinicalFinding.valueset"]     attributes --&gt; genericAttrib["generic.attrib"]     response --&gt; genericContent[generic.content]     genericContent --&gt; specify[specify]     genericContent --&gt; label[label]     genericContent --&gt; comment[comment]     genericContent --&gt; other["0..∞ #other"] </pre>																				
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre> &lt;response xml:id="" value=""&gt;   &lt;obbr:specify{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt; </pre>																				
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QName	Type	Fixed	Default	Use																	
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value	obbr:history.clinicalFinding.valueset			required																	
xml:id	xs:ID			optional																	
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="history.clinicalFinding.valueset" use="required"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

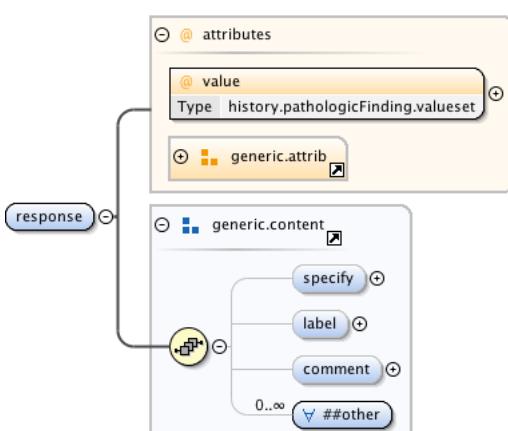
**Element history.pathologicFinding.item / pathologicFinding**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																																				
Properties	content: complex																																			
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																																			
Children	obbr:comment, obbr:label, obbr:specify, response																																			
Instance	<pathologicFinding date="" duration="" end="" xml:id="" start=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </pathologicFinding>																																			
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end	obbr:date.valueset			optional																																
start	obbr:date.valueset			optional																																
xml:id	xs:ID			optional																																
Source	<pre>&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="history.pathologicFinding.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="date"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>																																			
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																																			

### Element history.pathologicFinding.response / response

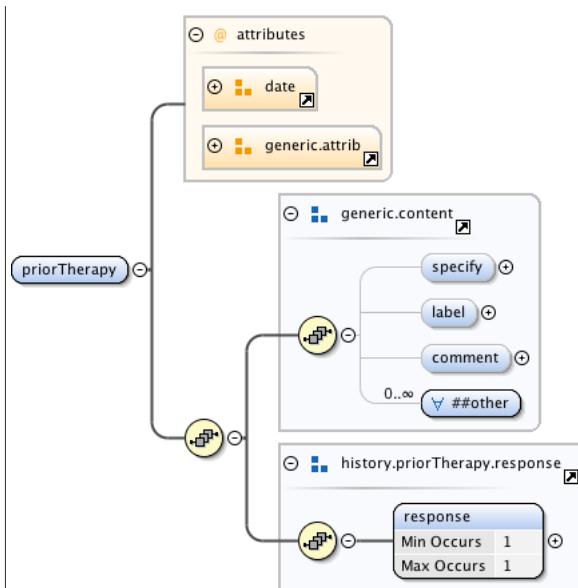
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<response xml:id="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </response>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:history.pathologicFinding.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:history.pathologicFinding.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
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<b>value</b>	obbr:history.pathologicFinding.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
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="history.pathologicFinding.valueset" use="required"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element history.priorTherapy.item / priorTherapy

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



Properties	content: complex																																			
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																																			
Children	obbr:comment, obbr:label, obbr:specify, response																																			
Instance	<priorTherapy date="" duration="" end="" xml:id="" start="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </priorTherapy>																																			
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>date</td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>duration</td> <td>xs:duration</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>end</td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>start</td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					date	obbr:date.valueset			optional	duration	xs:duration			optional	end	obbr:date.valueset			optional	start	obbr:date.valueset			optional	xml:id	xs:ID			optional
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end	obbr:date.valueset			optional																																
start	obbr:date.valueset			optional																																
xml:id	xs:ID			optional																																
Source	<x:element name="priorTherapy">   <x:complexType>     <x:sequence>       <x:group ref="generic.content"/>       <x:group ref="history.priorTherapy.response"/>     </x:sequence>     <x:attributeGroup ref="date"/>     <x:attributeGroup ref="generic.attrib"/>   </x:complexType> </x:element>																																			
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																																			

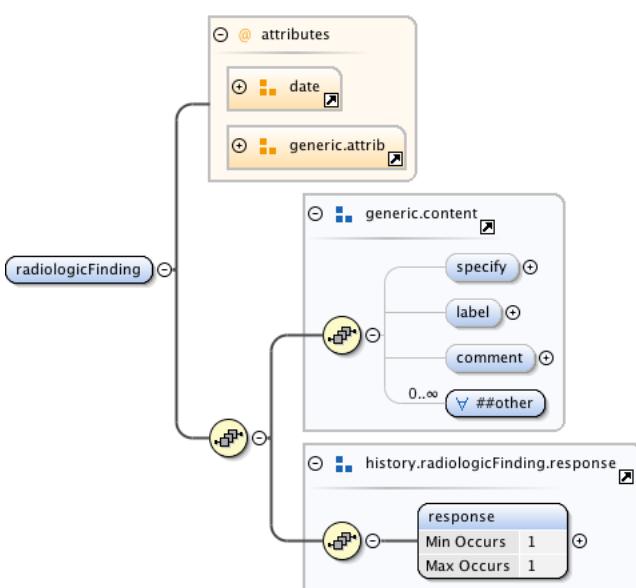
**Element history.priorTherapy.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
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QName	Type	Fixed	Default	Use																	
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<b>value</b>	obbr:history.priorTherapy.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;x:element name="response"&gt;   &lt;x:complexType&gt;     &lt;x:group ref="generic.content"/&gt;     &lt;x:attribute name="value" type="history.priorTherapy.valueset" use="required"/&gt;     &lt;x:attributeGroup ref="generic.attrib"/&gt;   &lt;/x:complexType&gt; &lt;/x:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

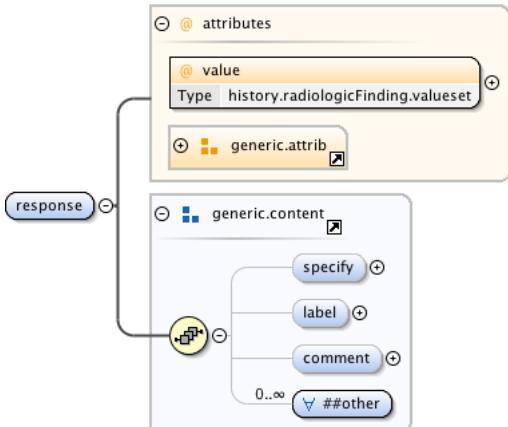
### Element history.radiologicFinding.item / radiologicFinding

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																																				
Properties	content: complex																																			
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																																			
Children	obbr:comment, obbr:label, obbr:specify, response																																			
Instance	<pre>&lt;radiologicFinding date="" duration="" end="" xml:id="" start=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/radiologicFinding&gt;</pre>																																			
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QName	Type	Fixed	Default	Use																																
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end	obbr:date.valueset			optional																																
start	obbr:date.valueset			optional																																
xml:id	xs:ID			optional																																
Source	<pre>&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="history.radiologicFinding.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="date"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>																																			
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																																			

### Element history.radiologicFinding.response / response

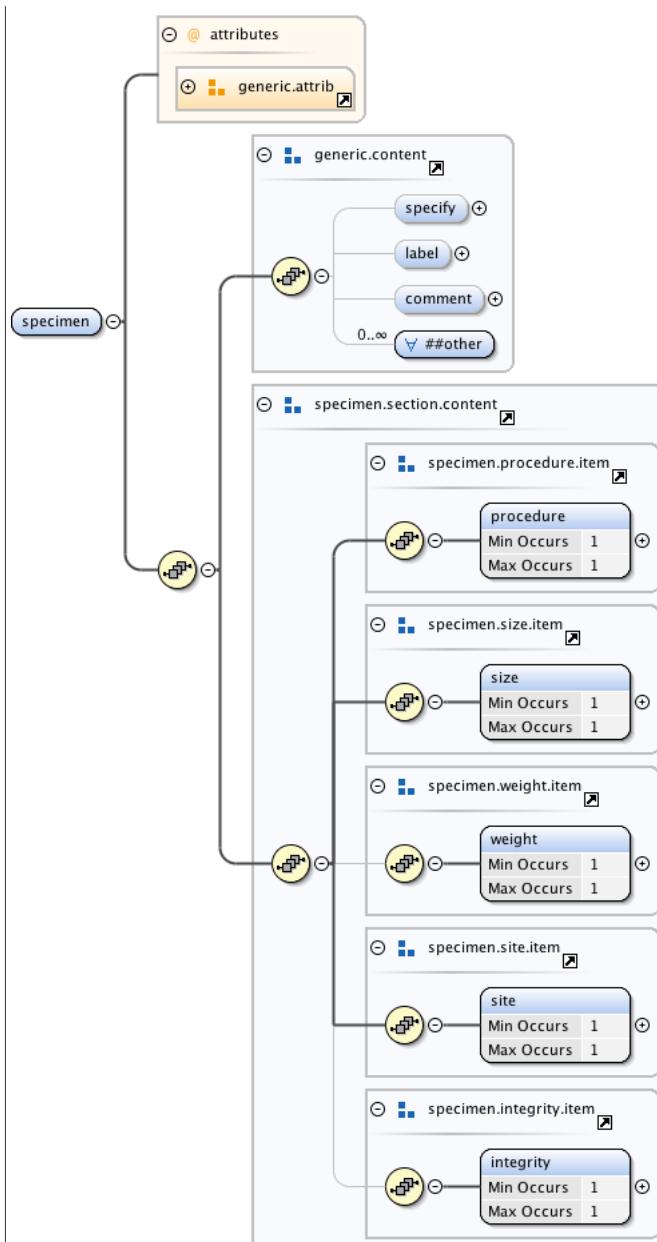
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:history.radiologicFinding.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:history.radiologicFinding.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:history.radiologicFinding.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;x:element name="response"&gt;   &lt;x:complexType&gt;     &lt;x:group ref="generic.content"/&gt;     &lt;x:attribute name="value" type="history.radiologicFinding.valueset" use="required"/&gt;     &lt;x:attributeGroup ref="generic.attrib"/&gt;   &lt;/x:complexType&gt; &lt;/x:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element specimen.section / specimen

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



Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , procedure , size , weight , site , integrity															
Children	integrity, obbr:comment, obbr:label, obbr:specify, procedure, site, size, weight															
Instance	<pre> &lt;specimen xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/specimen&gt;   </pre>															
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QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																
<b>xml:id</b>	xs:ID			optional												
Source	<pre> &lt;xss:element name="specimen"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       </pre>															

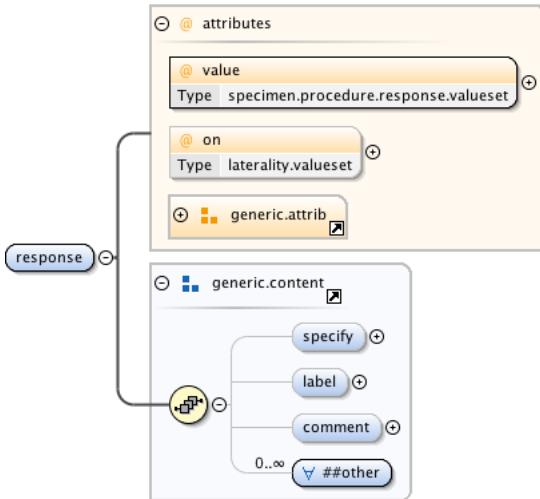
	<pre> &lt;xs:group ref="generic.content"/&gt; &lt;xs:group ref="specimen.section.content"/&gt; &lt;/xs:sequence&gt; &lt;xs:attributeGroup ref="generic.attrib"/&gt; &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

## Element specimen.procedure.item / procedure

Namespace	http://purl.org/obbr/pathology/cancer/															
Diagram	<pre> classDiagram     class procedure {         @ attributes         + generic.attrib     }     class generic.content {         + specify         + label         + comment         0..∞ ##other     }     class specimen.procedure.response {         + response         Min Occurs: 1         Max Occurs: unbounded     }     procedure "1..1" -- "1..∞" generic.content     generic.content "*" -- "1..∞" specimen.procedure.response </pre>															
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response+															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre> &lt;procedure xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/procedure&gt; </pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																
xml:id	xs:ID			optional												
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.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

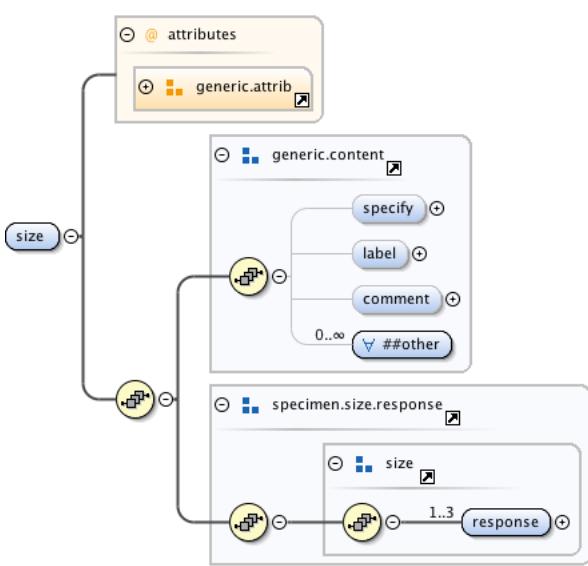
## Element specimen.procedure.response / response

Namespace	http://purl.org/obbr/pathology/cancer/
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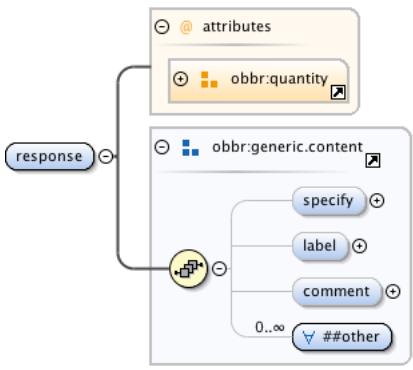
Diagram																										
Properties	<p>content: complex</p> <p>maxOccurs: unbounded</p>																									
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																									
Children	obbr:comment, obbr:label, obbr:specify																									
Instance	<pre>&lt;response xml:id="" on="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																									
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td><td></td><td></td><td></td><td></td></tr> <tr> <td>on</td><td>obbr:laterality.valueset</td><td></td><td></td><td>optional</td></tr> <tr> <td>value</td><td>obbr:specimen.procedure.response.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td>xml:id</td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					on	obbr:laterality.valueset			optional	value	obbr:specimen.procedure.response.valueset			required	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																						
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																										
on	obbr:laterality.valueset			optional																						
value	obbr:specimen.procedure.response.valueset			required																						
xml:id	xs:ID			optional																						
Source	<pre>&lt;xss:element maxOccurs="unbounded" name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="specimen.procedure.response.valueset" use="required"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																									
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																									

**Element specimen.size.item / size**

Namespace	http://purl.org/obbr/pathology/cancer/
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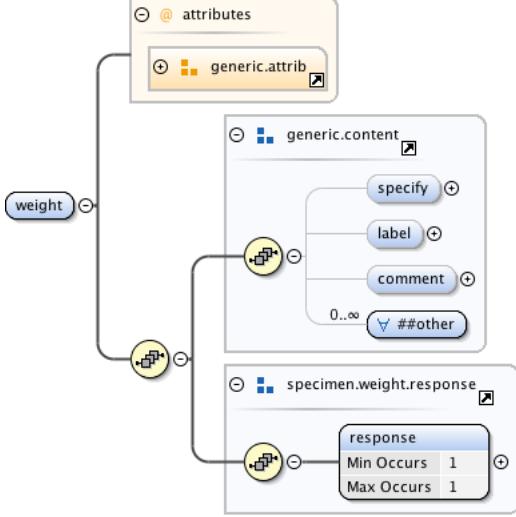
Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , obbr:response{1,3}															
Children	obbr:comment, obbr:label, obbr:response, obbr:specify															
Instance	<pre>&lt;size xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/size&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use												
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																
<b>xml:id</b>	xs:ID			optional												
Source	<pre>&lt;xs:element name="size"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="specimen.size.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element obbr:size / obbr:response

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

Properties	content: complex minOccurs: 1 maxOccurs: 3																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<obbr:response relation="" unit="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </obbr:response>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>relation</b></td><td>obbr:relation.valueset</td><td></td><td></td><td>optional</td></tr> <tr> <td><b>unit</b></td><td>obbr:lengthUnit.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>value</b></td><td>obbr:scalar.valueset</td><td></td><td></td><td>required</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>relation</b>	obbr:relation.valueset			optional	<b>unit</b>	obbr:lengthUnit.valueset			required	<b>value</b>	obbr:scalar.valueset			required
QName	Type	Fixed	Default	Use																	
<b>relation</b>	obbr:relation.valueset			optional																	
<b>unit</b>	obbr:lengthUnit.valueset			required																	
<b>value</b>	obbr:scalar.valueset			required																	
Source	<xs:element maxOccurs="3" minOccurs="1" name="response"> <xs:complexType> <xs:group ref="obbr:generic.content"/> <xs:attributeGroup ref="obbr:quantity"/> </xs:complexType> </xs:element>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd																				

**Element specimen.weight.item / weight**

Namespace	http://purl.org/obbr/pathology/cancer/															
Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<weight xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </weight>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use												
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																
<b>xml:id</b>	xs:ID			optional												

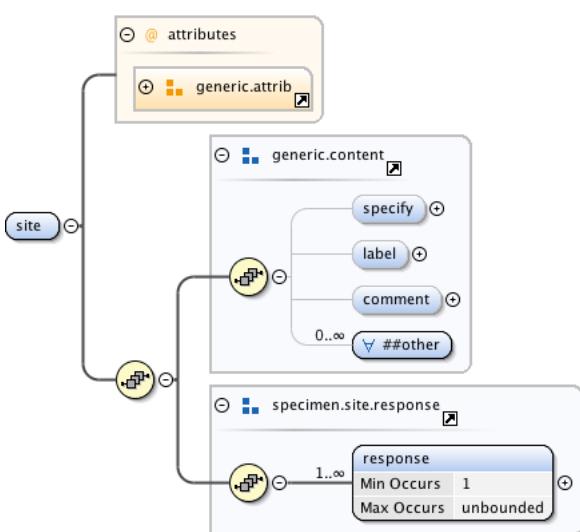
Source	<pre>&lt;xss:element name="weight"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="specimen.weight.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element specimen.weight.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/																														
Diagram	<pre> classDiagram     class response {         @ attributes         + weight         + generic.attrib         generic.content {             + specify             + label             + comment             0..∞ #other         }     }   </pre>																														
Properties	content: complex																														
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																														
Children	obbr:comment, obbr:label, obbr:specify																														
Instance	<pre>&lt;response xml:id="" relation="" unit="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																														
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>relation</td> <td>obbr:relation.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>unit</td> <td>obbr:weightUnit.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>value</td> <td>obbr:scalar.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					relation	obbr:relation.valueset			optional	unit	obbr:weightUnit.valueset			required	value	obbr:scalar.valueset			required	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																											
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																															
relation	obbr:relation.valueset			optional																											
unit	obbr:weightUnit.valueset			required																											
value	obbr:scalar.valueset			required																											
xml:id	xs:ID			optional																											
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attributeGroup ref="weight"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																														
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																														

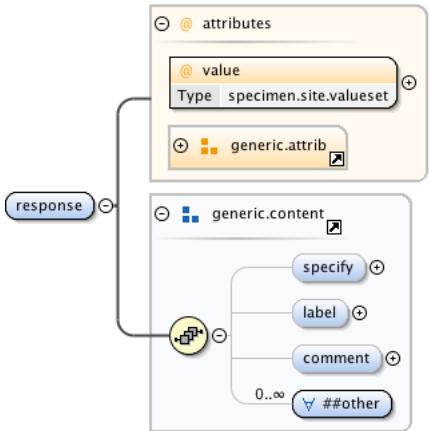
**Element specimen.site.item / site**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response+															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<site xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </site>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre> &lt;xss:element name="site"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="specimen.site.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt; </pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

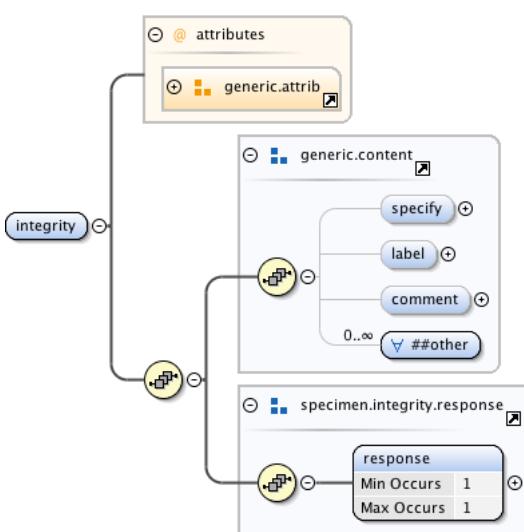
### Element specimen.site.response / response

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

Diagram																					
Properties	<p>content: complex</p> <p>maxOccurs: unbounded</p>																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>value</b></td> <td>obbr:specimen.site.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:specimen.site.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:specimen.site.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
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.site.valueset" use="required"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

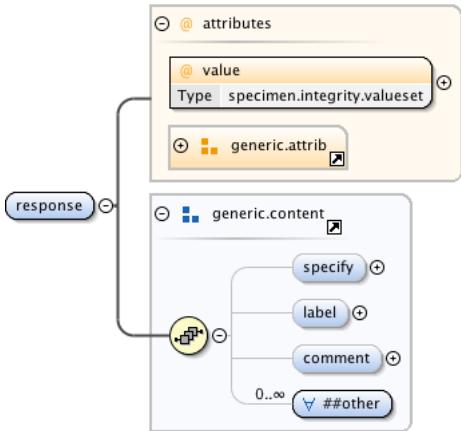
## Element specimen.integrity.item / integrity

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre>&lt;integrity xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/integrity&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="integrity"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="specimen.integrity.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element specimen.integrity.response / response

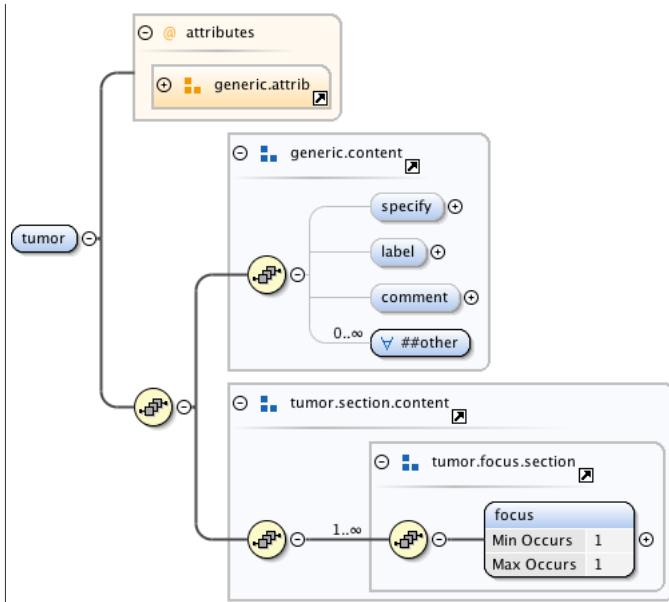
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:specimen.integrity.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:specimen.integrity.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:specimen.integrity.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="specimen.integrity.valueset" use="required"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

## Element tumor.section / tumor

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

## Diagram



## Properties

content: complex

## Model

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

## Children

focus, obbr:comment, obbr:label, obbr:specify

## Instance

```

<tumor xml:id="">
  <obbr:specify>{0,1}</obbr:specify>
  <obbr:label>{0,1}</obbr:label>
  <obbr:comment>{0,1}</obbr:comment>
</tumor>
  
```

## Attributes

QName	Type	Fixed	Default	Use
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'				
xml:id	xs:ID			optional

## Source

```

<xs:element name="tumor">
  <xs:complexType>
    <xs:sequence>
      <xs:group ref="generic.content"/>
      <xs:group ref="tumor.section.content"/>
    </xs:sequence>
    <xs:attributeGroup ref="generic.attrib"/>
  </xs:complexType>
</xs:element>
  
```

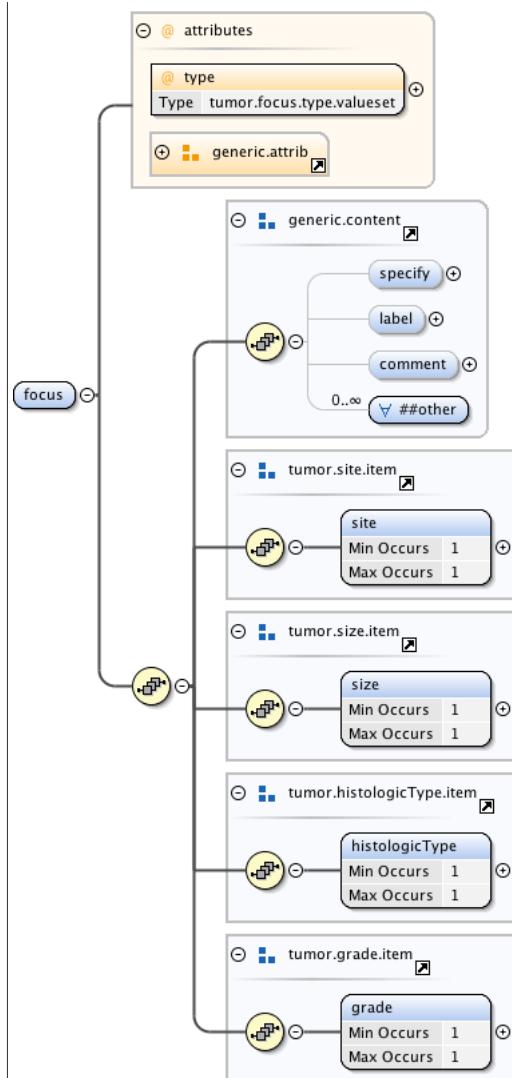
## Schema location

file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element tumor.focus.section / focus**

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



Properties	content: complex				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , site , size , histologicType , grade				
Children	grade, histologicType, obbr:comment, obbr:label, obbr:specify, site, size				
Instance	<focus xml:id="" type=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </focus>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	type	obbr:tumor.focus.type.valueset			required
	xml:id	xs:ID			optional
Source	<xss:element name="focus"> <xss:complexType> <xss:sequence> <xss:group ref="generic.content"/> <xss:group ref="tumor.size.item"/> <xss:group ref="tumor.size.item"/> <xss:group ref="tumor.histologicType.item"/> <xss:group ref="tumor.grade.item"/>				

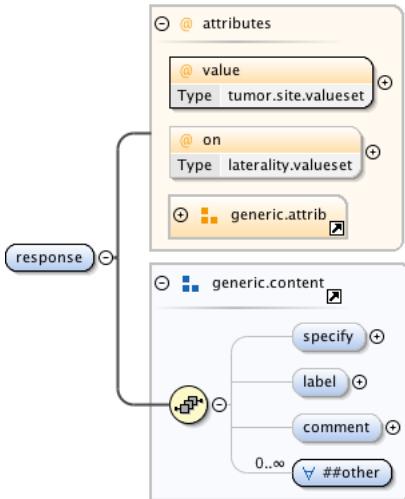
	<pre> &lt;/xs:sequence&gt; &lt;xs:attribute name="type" type="tumor.focus.type.valueset" use="required"/&gt; &lt;xss:attributeGroup ref="generic.attrib"/&gt; &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element tumor.site.item / site**

Namespace	http://purl.org/obbr/pathology/cancer/															
Diagram	<pre> classDiagram     class site {         @ attributes         + generic.attrib         + generic.content {             + specify             + label             + comment             + 0..∞ #other         }         + tumor.site.response {             + response             Min Occurs 1             Max Occurs 1         }     } </pre>															
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre> &lt;site xml:id=""&gt;     &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;     &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;     &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/site&gt; </pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																
xml:id	xs:ID			optional												
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="tumor.site.response"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="generic.attrib"/&gt;     &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

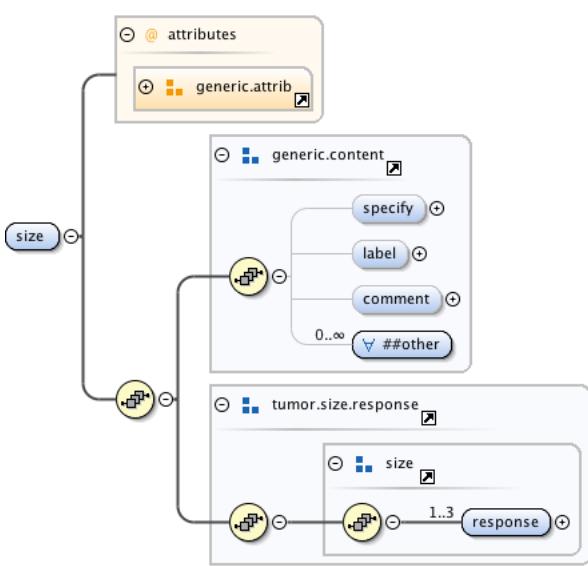
**Element tumor.site.response / response**

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

Diagram																										
Properties	content: complex																									
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																									
Children	obbr:comment, obbr:label, obbr:specify																									
Instance	<response xml:id="" on="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </response>																									
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>on</b></td><td>obbr:laterality.valueset</td><td></td><td></td><td>optional</td></tr> <tr> <td><b>value</b></td><td>obbr:tumor.site.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>on</b>	obbr:laterality.valueset			optional	<b>value</b>	obbr:tumor.site.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																						
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																										
<b>on</b>	obbr:laterality.valueset			optional																						
<b>value</b>	obbr:tumor.site.valueset			required																						
<b>xml:id</b>	xs:ID			optional																						
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="tumor.site.valueset" use="required"/&gt;     &lt;xss:attribute name="on" type="laterality.valueset"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																									
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																									

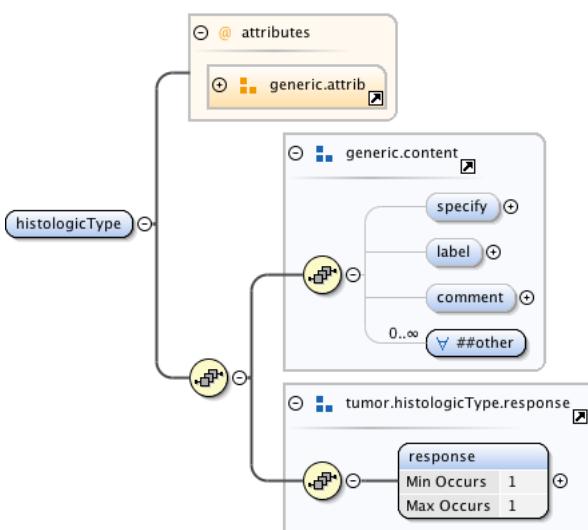
### Element tumor.size.item / size

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , obbr:response{1,3}															
Children	obbr:comment, obbr:label, obbr:response, obbr:specify															
Instance	<pre>&lt;size xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/size&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>					<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use												
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>																
<b>xml:id</b>	xs:ID			optional												
Source	<pre>&lt;xs:element name="size"&gt;   &lt;xs:complexType&gt;     &lt;xs:sequence&gt;       &lt;xs:group ref="generic.content"/&gt;       &lt;xs:group ref="tumor.size.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

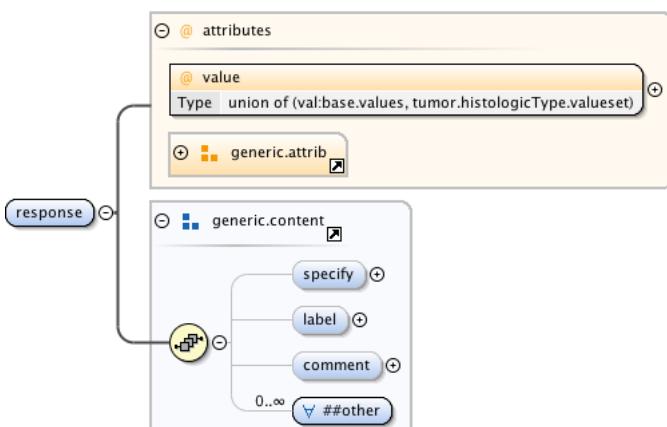
### Element tumor.histologicType.item / histologicType

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

Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre>&lt;histologicType xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/histologicType&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>					<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use												
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>																
<b>xml:id</b>	xs:ID			optional												
Source	<pre>&lt;x:element name="histologicType"&gt;   &lt;x:complexType&gt;     &lt;x:sequence&gt;       &lt;x:group ref="generic.content"/&gt;       &lt;x:group ref="tumor.histologicType.response"/&gt;     &lt;/x:sequence&gt;     &lt;x:attributeGroup ref="generic.attrib"/&gt;   &lt;/x:complexType&gt; &lt;/x:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

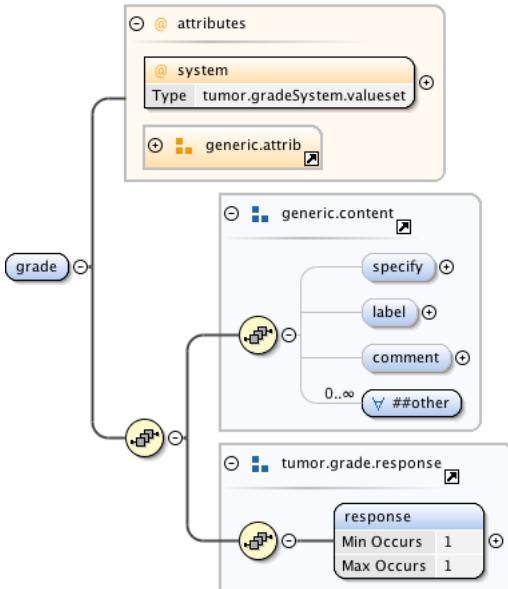
### Element tumor.histologicType.response / response

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

Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
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QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	union of(obbr:base.values, obbr:tumor.histologicType.valueset)			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" use="required"&gt;       &lt;xss:simpleType&gt;         &lt;xss:union memberTypes="val:base.values tumor.histologicType.valueset"/&gt;       &lt;/xss:simpleType&gt;     &lt;/xss:attribute&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

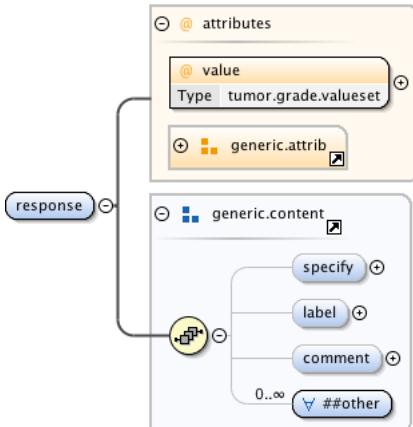
### Element tumor.grade.item / grade

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

Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																				
Children	obbr:comment, obbr:label, obbr:specify, response																				
Instance	<pre>&lt;grade xml:id="" system=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/grade&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>system</td> <td>obbr:tumor.gradeSystem.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					system	obbr:tumor.gradeSystem.valueset			required	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																	
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																					
system	obbr:tumor.gradeSystem.valueset			required																	
xml:id	xs:ID			optional																	
Source	<pre>&lt;xss:element name="grade"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="tumor.grade.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attribute name="system" type="tumor.gradeSystem.valueset" use="required"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

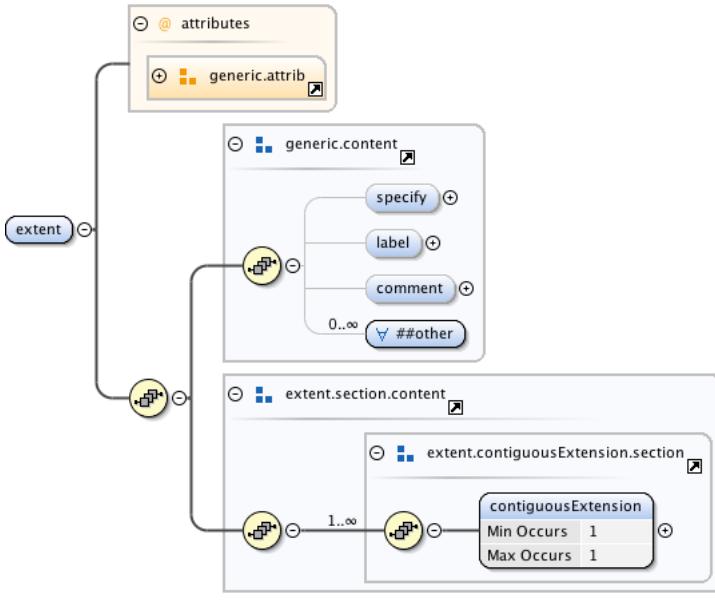
### Element tumor.grade.response / response

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

Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>value</b></td> <td>obbr:tumor.grade.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:tumor.grade.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:tumor.grade.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xsd:element name="response"&gt;   &lt;xsd:complexType&gt;     &lt;xsd:group ref="generic.content"/&gt;     &lt;xsd:attribute name="value" type="tumor.grade.valueset" use="required"/&gt;     &lt;xsd:attributeGroup ref="generic.attrib"/&gt;   &lt;/xsd:complexType&gt; &lt;/xsd:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element extent.section / extent

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , contiguousExtension															
Children	contiguousExtension, obbr:comment, obbr:label, obbr:specify															
Instance	<pre>&lt;extent xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/extent&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="extent"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="extent.section.content"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element extent.contiguousExtension.section / contiguousExtension

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status															
Children	obbr:comment, obbr:label, obbr:specify, status															
Instance	<contiguousExtension xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </contiguousExtension>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td><td></td><td></td><td></td><td></td></tr> <tr> <td>xml:id</td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="contiguousExtension"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group maxOccurs="unbounded" minOccurs="0"       ref="extent.contiguousExtension.status.section"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

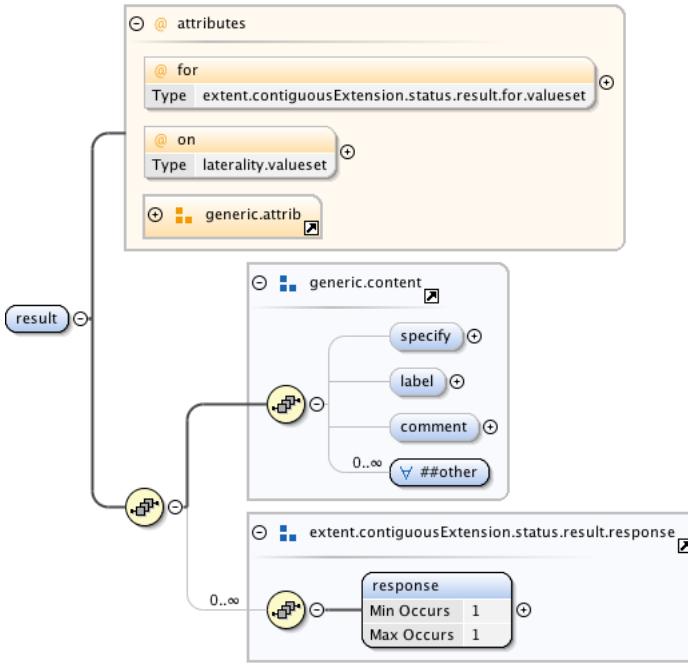
### Element extent.contiguousExtension.status.section / status

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , result , distance																				
Children	distance, obbr:comment, obbr:label, obbr:specify, result																				
Instance	<pre>&lt;status xml:id="" of=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/status&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>of</td> <td>obbr:extent.status.of.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					of	obbr:extent.status.of.valueset			optional	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																	
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																					
of	obbr:extent.status.of.valueset			optional																	
xml:id	xs:ID			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 maxOccurs="2" ref="extent.contiguousExtension.status.result.item"/&gt;       &lt;xs:group maxOccurs="2" minOccurs="0" ref="extent.contiguousExtension.status.distance.item"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="of" type="extent.status.of.valueset"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

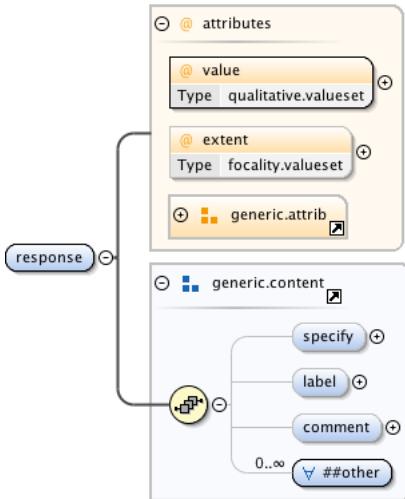
### Element extent.contiguousExtension.status.result.item / result

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																										
Properties	content: complex																									
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																									
Children	obbr:comment, obbr:label, obbr:specify, response																									
Instance	<result for="" xml:id="" on=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </result>																									
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>for</b></td> <td>obbr:extent.contiguousExtension.status.result.for.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>on</b></td> <td>obbr:laterality.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>for</b>	obbr:extent.contiguousExtension.status.result.for.valueset			optional	<b>on</b>	obbr:laterality.valueset			optional	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																						
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																										
<b>for</b>	obbr:extent.contiguousExtension.status.result.for.valueset			optional																						
<b>on</b>	obbr:laterality.valueset			optional																						
<b>xml:id</b>	xs:ID			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="extent.contiguousExtension.status.result.response"/&gt;     &lt;/xs:sequence&gt;     &lt;xs:attribute name="for" type="extent.contiguousExtension.status.result.for.valueset"/&gt;     &lt;xs:attribute name="on" type="laterality.valueset"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>																									
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																									

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

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																										
Properties	content: complex																									
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																									
Children	obbr:comment, obbr:label, obbr:specify																									
Instance	<pre>&lt;response extent="" xml:id="" value=""&gt;   &lt;obbr:specify{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																									
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td><td></td><td></td><td></td><td></td></tr> <tr> <td>extent</td><td>obbr:focality.valueset</td><td></td><td></td><td>optional</td></tr> <tr> <td>value</td><td>obbr:qualitative.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td>xml:id</td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					extent	obbr:focality.valueset			optional	value	obbr:qualitative.valueset			required	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																						
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																										
extent	obbr:focality.valueset			optional																						
value	obbr:qualitative.valueset			required																						
xml:id	xs:ID			optional																						
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="qualitative.valueset" use="required"/&gt;     &lt;xss:attribute name="extent" type="focality.valueset"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																									
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																									

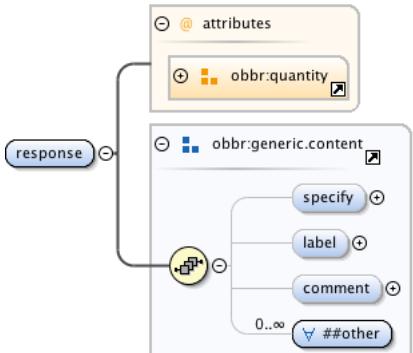
**Element extent.contiguousExtension.status.distance.item / distance**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																										
Properties	content: complex																									
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , obbr:response																									
Children	obbr:comment, obbr:label, obbr:response, obbr:specify																									
Instance	<distance for="" from="" xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </distance>																									
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>for</b></td> <td>obbr:extent.contiguousExtension.status.distance.for.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>from</b></td> <td>obbr:base.values</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>for</b>	obbr:extent.contiguousExtension.status.distance.for.valueset			optional	<b>from</b>	obbr:base.values			optional	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																						
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																										
<b>for</b>	obbr:extent.contiguousExtension.status.distance.for.valueset			optional																						
<b>from</b>	obbr:base.values			optional																						
<b>xml:id</b>	xs:ID			optional																						
Source	<pre> &lt;xss:element name="distance"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="length"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attribute name="for"       type="extents.contiguousExtension.status.distance.for.valueset"/&gt;     &lt;xss:attribute name="from" type="val:base.values"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt; </pre>																									
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																									

### Element obbr:length / obbr:response

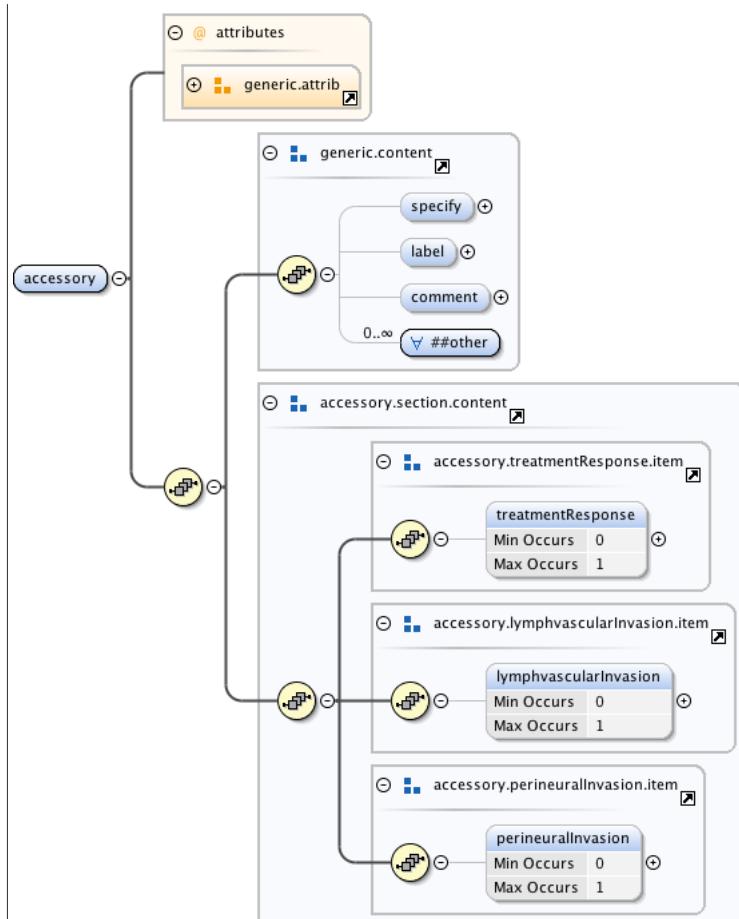
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<obbr:response relation="" unit="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </obbr:response>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>relation</b></td> <td>obbr:relation.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>unit</b></td> <td>obbr:lengthUnit.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td><b>value</b></td> <td>obbr:scalar.valueset</td> <td></td> <td></td> <td>required</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>relation</b>	obbr:relation.valueset			optional	<b>unit</b>	obbr:lengthUnit.valueset			required	<b>value</b>	obbr:scalar.valueset			required
QName	Type	Fixed	Default	Use																	
<b>relation</b>	obbr:relation.valueset			optional																	
<b>unit</b>	obbr:lengthUnit.valueset			required																	
<b>value</b>	obbr:scalar.valueset			required																	
Source	<pre>&lt;xsd:element name="response"&gt;   &lt;xsd:complexType&gt;     &lt;xsd:group ref="obbr:generic.content"/&gt;     &lt;xsd:attributeGroup ref="obbr:quantity"/&gt;   &lt;/xsd:complexType&gt; &lt;/xsd:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd																				

### Element accessory.section / accessory

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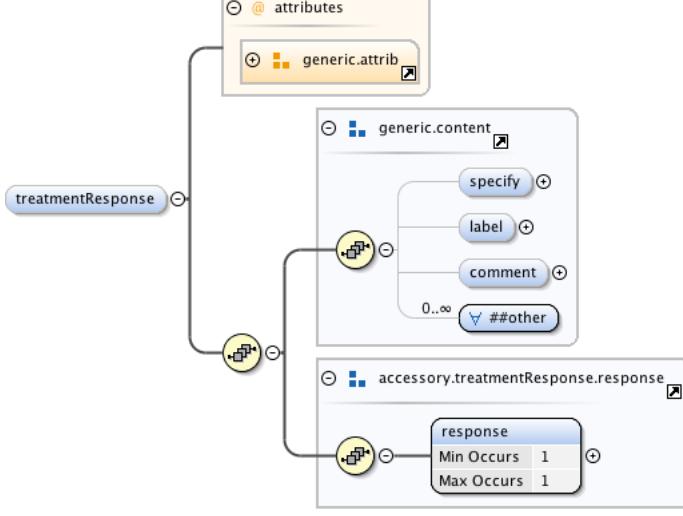
## Diagram



Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , treatmentResponse{0,1} , lymphvascularInvasion{0,1} , perineuralInvasion{0,1}															
Children	lymphvascularInvasion, obbr:comment, obbr:label, obbr:specify, perineuralInvasion, treatmentResponse															
Instance	<pre> &lt;accessory xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/accessory&gt; </pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use												
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																
<b>xml:id</b>	xs:ID			optional												
Source	<pre> &lt;xss:element name="accessory"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="accessory.section.content"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt; </pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

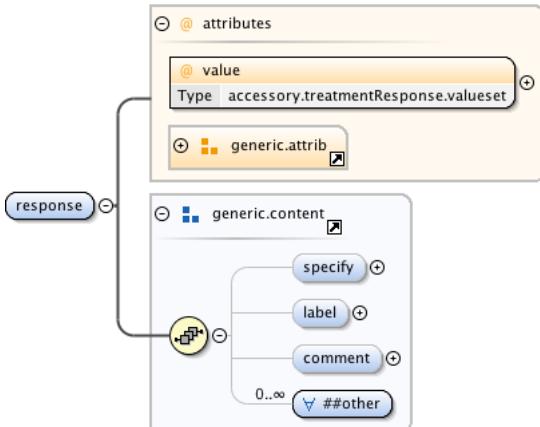
**Element accessory.treatmentResponse.item / treatmentResponse**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	<p>content: complex</p> <p>minOccurs: 0</p>															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre>&lt;treatmentResponse xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/treatmentResponse&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element minOccurs="0" name="treatmentResponse"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="accessory.treatmentResponse.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

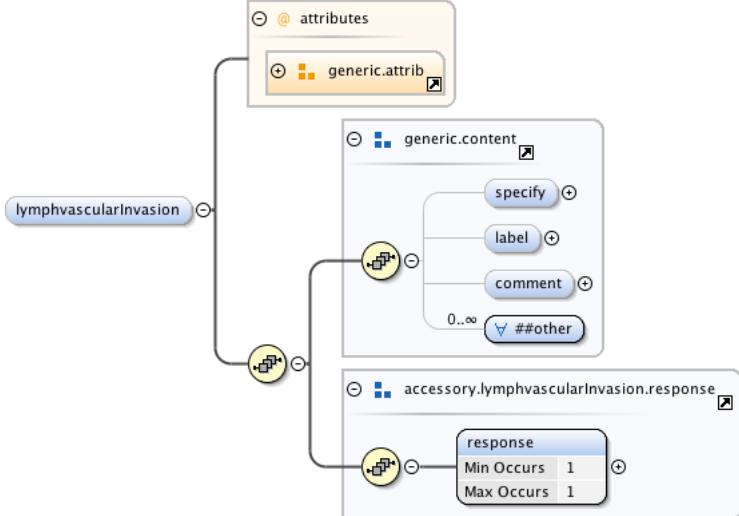
### Element accessory.treatmentResponse.response / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:accessory.treatmentResponse.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:accessory.treatmentResponse.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:accessory.treatmentResponse.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="accessory.treatmentResponse.valueset" use="required"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

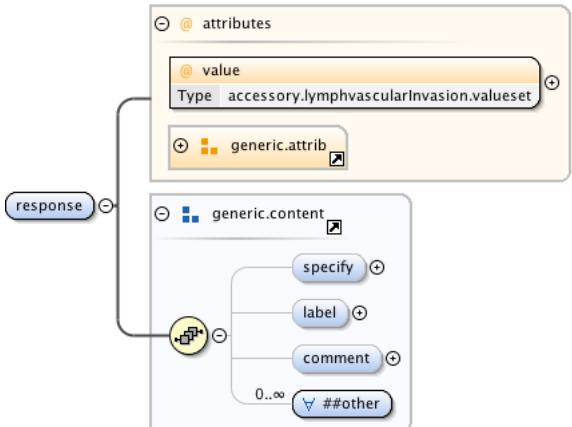
### Element accessory.lymphvascularInvasion.item / lymphvascularInvasion

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	<p>content: complex</p> <p>minOccurs: 0</p>															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre>&lt;lymphvascularInvasion xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/lymphvascularInvasion&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'</td><td></td><td></td><td></td><td></td></tr> <tr> <td>xml:id</td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element minOccurs="0" name="lymphvascularInvasion"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="accessory.lymphvascularInvasion.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

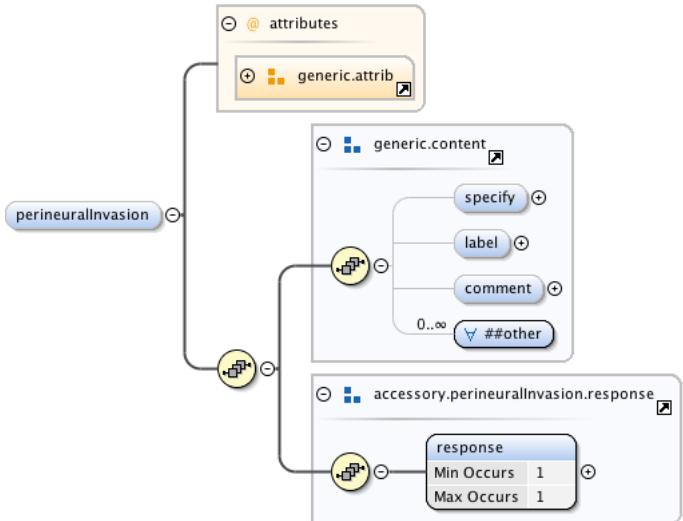
### Element accessory.lymphvascularInvasion.response / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>value</b></td> <td>obbr:accessory.lymphvascularInvasion.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:accessory.lymphvascularInvasion.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:accessory.lymphvascularInvasion.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="accessory.lymphvascularInvasion.valueset" use="required"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element accessory.perineuralInvasion.item / perineuralInvasion

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	<p>content: complex</p> <p>minOccurs: 0</p>															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre>&lt;perineuralInvasion xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/perineuralInvasion&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element minOccurs="0" name="perineuralInvasion"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="accessory.perineuralInvasion.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

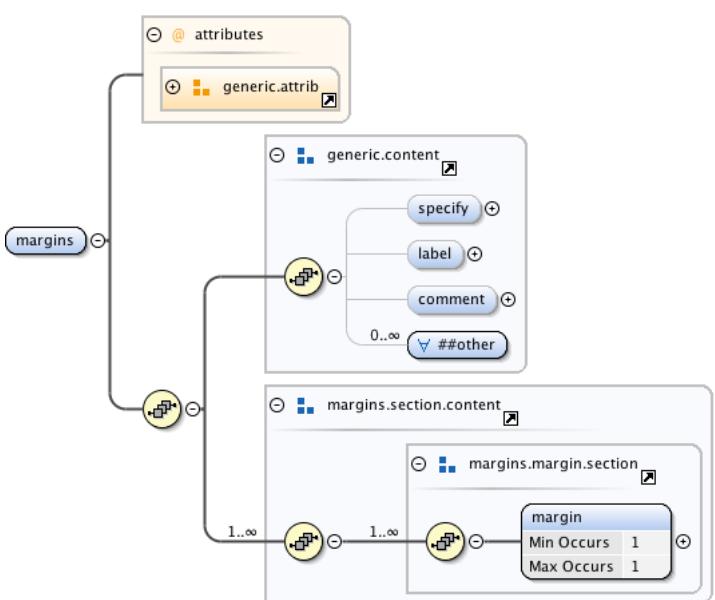
### Element accessory.perineuralInvasion.response / response

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

Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:accessory.perineuralInvasion.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:accessory.perineuralInvasion.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:accessory.perineuralInvasion.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xsd:element name="response"&gt;   &lt;xsd:complexType&gt;     &lt;xsd:group ref="generic.content"/&gt;     &lt;xsd:attribute name="value" type="accessory.perineuralInvasion.valueset"       use="required"/&gt;     &lt;xsd:attributeGroup ref="generic.attrib"/&gt;   &lt;/xsd:complexType&gt; &lt;/xsd:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element margins.section / margins

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer' , margin															
Children	margin, obbr:comment, obbr:label, obbr:specify															
Instance	<Margins xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </Margins>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="margins"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group maxOccurs="unbounded" ref="margins.section.content"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element margins.margin.section / margin

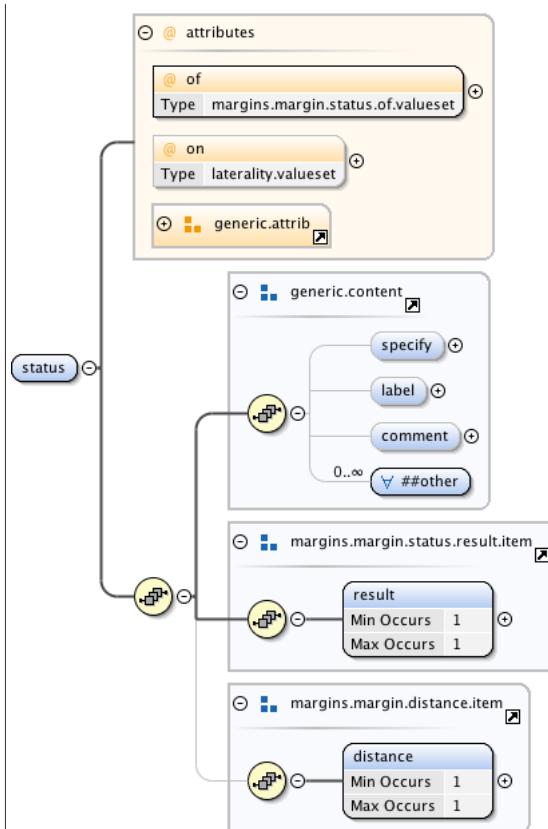
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status															
Children	obbr:comment, obbr:label, obbr:specify, status															
Instance	<margin xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </margin>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="margin"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="margins.margin.section.content"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element margins.margin.status.section / status

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



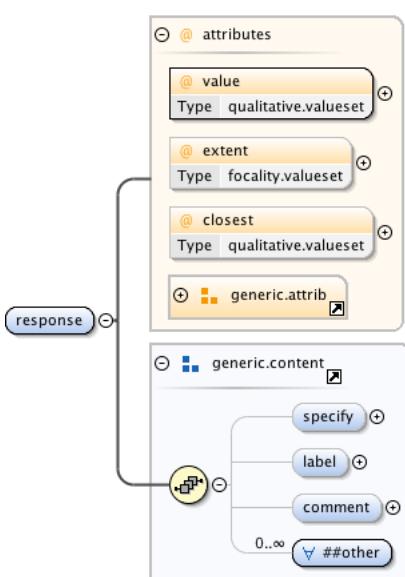
Properties	content: complex																									
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , result , distance																									
Children	distance, obbr:comment, obbr:label, obbr:specify, result																									
Instance	<status xml:id="" of="" on="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </status>																									
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>of</td> <td>obbr:margins.margin.status.of.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>on</td> <td>obbr:laterality.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					of	obbr:margins.margin.status.of.valueset			required	on	obbr:laterality.valueset			optional	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																						
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																										
of	obbr:margins.margin.status.of.valueset			required																						
on	obbr:laterality.valueset			optional																						
xml:id	xs:ID			optional																						
Source	<xs:element name="status">   <xs:complexType>     <xs:sequence>       <xs:group ref="generic.content"/>       <xs:group ref="margins.margin.status.result.item"/>       <xs:group minOccurs="0" ref="margins.margin.distance.item"/>     </xs:sequence>     <xs:attribute name="of" type="margins.margin.status.of.valueset" use="required"/>     <xs:attribute name="on" type="laterality.valueset"/>     <xs:attributeGroup ref="generic.attrib"/>   </xs:complexType> </xs:element>																									
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																									

**Element margins.margin.status.result.item / result**

Namespace	http://purl.org/obbr/pathology/cancer/																							
Diagram	<pre> classDiagram     class margins.margin.status.result.item {         @ for margins.margin.status.result.for.valueset         generic.attrib         generic.content {             specify             label             comment             #other         }         margins.margin.status.result.response {             response             Min Occurs 1             Max Occurs 1         }     } </pre>																							
Properties	content: complex																							
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																							
Children	obbr:comment, obbr:label, obbr:specify, response																							
Instance	<result for="" xml:id="">     <obbr:specify>{0,1}</obbr:specify>     <obbr:label>{0,1}</obbr:label>     <obbr:comment>{0,1}</obbr:comment> </result>																							
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>for</td> <td>obbr:margins.margin.status.result.for.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>				QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					for	obbr:margins.margin.status.result.for.valueset			required	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																				
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																								
for	obbr:margins.margin.status.result.for.valueset			required																				
xml:id	xs:ID			optional																				
Source	<xss:element name="result">     <xss:complexType>         <xss:sequence>             <xss:group ref="generic.content"/>             <xss:group ref="margins.margin.status.result.response"/>         </xss:sequence>         <xss:attribute name="for" type="margins.margin.status.result.for.valueset" use="required"/>         <xss:attributeGroup ref="generic.attrib"/>     </xss:complexType> </xss:element>																							
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																							

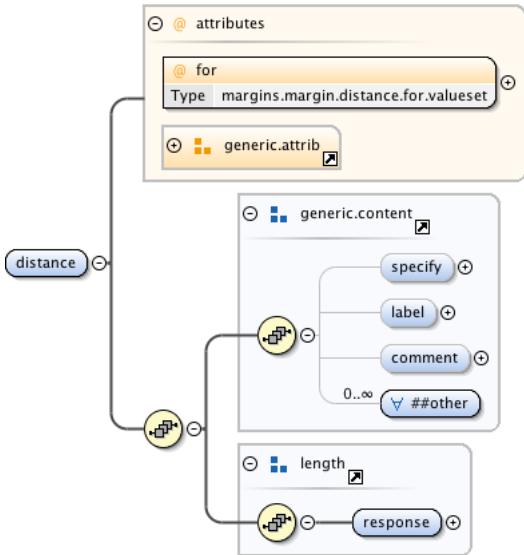
**Element margins.margin.status.result.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																															
Properties	content: complex																														
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																														
Children	obbr:comment, obbr:label, obbr:specify																														
Instance	<response closest="" extent="" xml:id="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </response>																														
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>closest</td> <td>obbr:qualitative.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>extent</td> <td>obbr:focality.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>value</td> <td>obbr:qualitative.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					closest	obbr:qualitative.valueset			optional	extent	obbr:focality.valueset			optional	value	obbr:qualitative.valueset			required	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																											
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																															
closest	obbr:qualitative.valueset			optional																											
extent	obbr:focality.valueset			optional																											
value	obbr:qualitative.valueset			required																											
xml:id	xs:ID			optional																											
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="qualitative.valueset" use="required"/&gt;     &lt;xss:attribute name="extent" type="focality.valueset"/&gt;     &lt;xss:attribute name="closest" type="qualitative.valueset"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																														
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																														

**Element margins.margin.distance.item / distance**

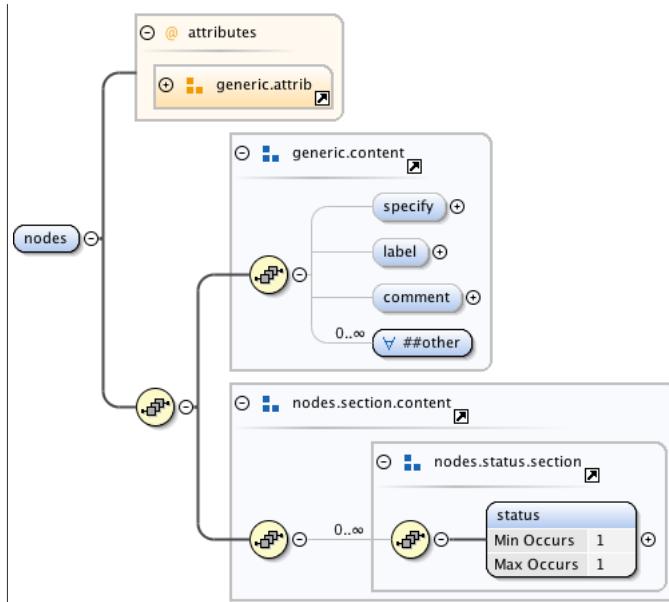
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , obbr:response																				
Children	obbr:comment, obbr:label, obbr:response, obbr:specify																				
Instance	<pre>&lt;distance for="" xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/distance&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>for</b></td><td>obbr:margins.margin.distance.for.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>for</b>	obbr:margins.margin.distance.for.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>for</b>	obbr:margins.margin.distance.for.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="distance"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="length"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attribute name="for" type="margins.margin.distance.for.valueset" use="required"/&gt;     &lt;!--&lt;xss:attribute name="from" type="margins.margin.distance.from.valueset"/&gt;--&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

## Element nodes.section / nodes

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

## Diagram



## Properties

content: complex

## Model

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

## Children

obbr:comment, obbr:label, obbr:specify, status

## Instance

```

<nodes xml:id="">
  <obbr:specify>{0,1}</obbr:specify>
  <obbr:label>{0,1}</obbr:label>
  <obbr:comment>{0,1}</obbr:comment>
</nodes>
  
```

## Attributes

QName	Type	Fixed	Default	Use
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'				
xml:id	xs:ID			optional

## Source

```

<xs:element name="nodes">
  <xs:complexType>
    <xs:sequence>
      <xs:group ref="generic.content"/>
      <xs:group ref="nodes.section.content"/>
    </xs:sequence>
    <xs:attributeGroup ref="generic.attrib"/>
  </xs:complexType>
</xs:element>
  
```

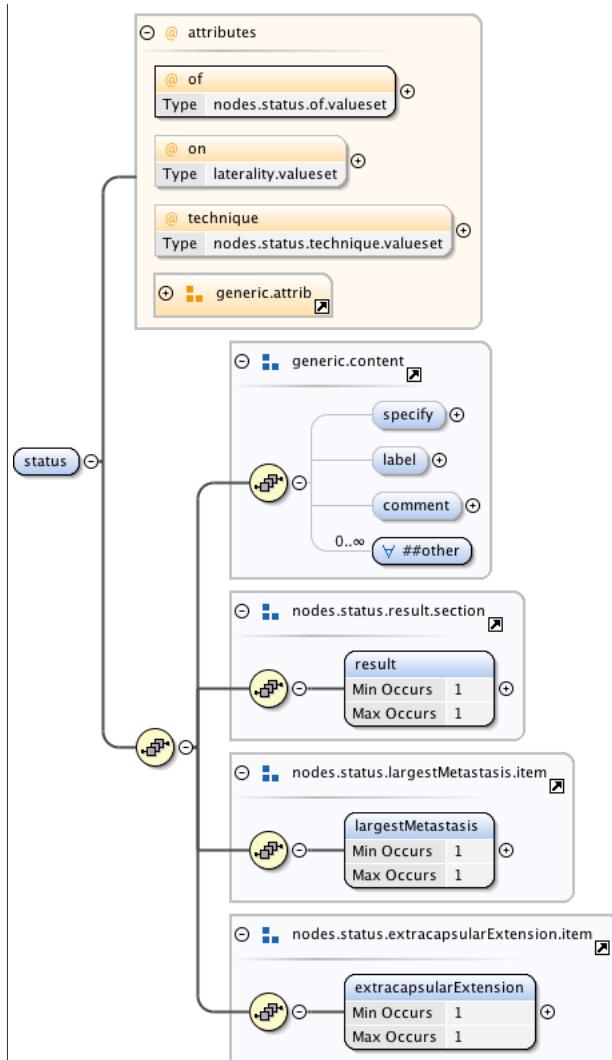
## Schema location

file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element nodes.status.section / status**

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

## Diagram



Properties	content: complex				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , result , largestMetastasis , extracapsularExtension				
Children	extracapsularExtension, largestMetastasis, obbr:comment, obbr:label, obbr:specify, result				
Instance	<status xml:id="" of="" on="" technique=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </status>				
Attributes	QName	Type	Fixed	Default	Use
	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
	of	obbr:nodes.status.of.valueset			required
	on	obbr:laterality.valueset			optional
	technique	obbr:nodes.status.technique.valueset			optional
	xml:id	xs:ID			optional
Source	<xss:element name="status"> <xss:complexType> <xss:sequence> <xss:group ref="generic.content"/> <xss:group ref="nodes.status.result.section"/> <xss:group ref="nodes.status.largestMetastasis.item"/>				

```

<xs:group ref="nodes.status.extracapsularExtension.item"/>
</xs:sequence>
<xs:attribute name="of" type="nodes.status.of.valueset" use="required"/>
<xs:attribute name="on" type="laterality.valueset"/>
<xs:attribute name="technique" type="nodes.status.technique.valueset"/>
<xs:attributeGroup ref="generic.attrib"/>
</xs:complexType>
</xs:element>

```

Schema location file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

## Element nodes.status.result.section / result

Namespace	http://purl.org/obbr/pathology/cancer/																									
Diagram	<pre> classDiagram     class result {         @ for Type nodes.status.result.for.valueset         @ type Type nodes.status.result.type.valueset         + generic.attrib         --&gt; generic.content         --&gt; nodes.status.result.response     }     class generic.content {         specify         label         comment         0..∞ #other     }     class nodes.status.result.response {         response         Min Occurs 1         Max Occurs 1     } </pre>																									
Properties	content: complex																									
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																									
Children	obbr:comment, obbr:label, obbr:specify, response																									
Instance	<result for="" xml:id="" type="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </result>																									
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>for</td> <td>obbr:nodes.status.result.for.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>type</td> <td>obbr:nodes.status.result.type.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'					for	obbr:nodes.status.result.for.valueset			required	type	obbr:nodes.status.result.type.valueset			optional	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																						
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'																										
for	obbr:nodes.status.result.for.valueset			required																						
type	obbr:nodes.status.result.type.valueset			optional																						
xml:id	xs:ID			optional																						
Source	<xs:element name="result">   <xs:complexType>     <xs:sequence>       <xs:group ref="generic.content"/>       <xs:group ref="nodes.status.result.response"/>     </xs:sequence>     <xs:attribute name="for" type="nodes.status.result.for.valueset" use="required"/>     <xs:attribute name="type" type="nodes.status.result.type.valueset"/>     <xs:attributeGroup ref="generic.attrib"/>   </xs:complexType> </xs:element>																									

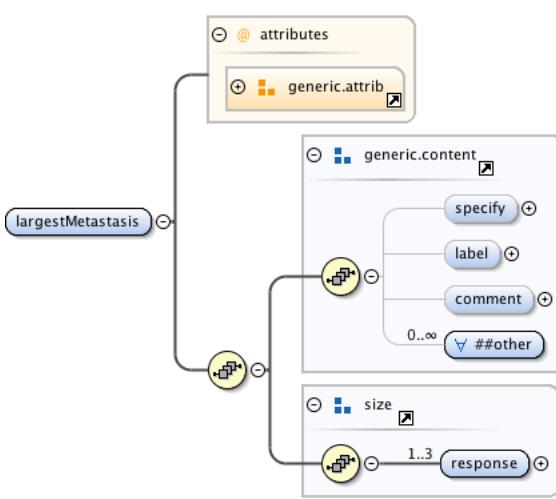
	</xs:element>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element nodes.status.result.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/																													
Diagram	<pre> classDiagram     class response {         @ attributes: tally, generic.attrib         generic.content {             specify             label             comment             0..∞ other         }     }   </pre>																													
Properties	content: complex																													
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																													
Children	obbr:comment, obbr:label, obbr:specify																													
Instance	<response xml:id="" relation="" value="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </response>																													
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>relation</td> <td>obbr:relation.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td>value</td> <td>obbr:tally.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>					QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					relation	obbr:relation.valueset			optional	value	obbr:tally.valueset			required	xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																										
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																														
relation	obbr:relation.valueset			optional																										
value	obbr:tally.valueset			required																										
xml:id	xs:ID			optional																										
Source	<pre> &lt;xs:element name="response"&gt;   &lt;xs:complexType&gt;     &lt;xs:group ref="generic.content"/&gt;     &lt;xs:attributeGroup ref="tally"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt; </pre>																													
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																													

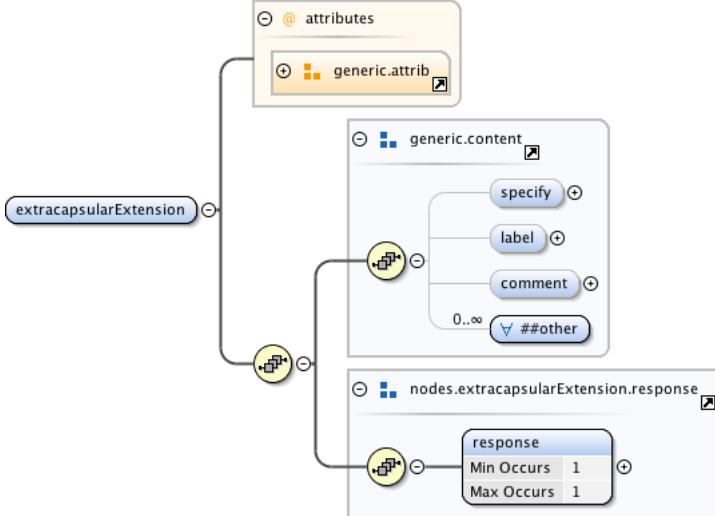
**Element nodes.status.largestMetastasis.item / largestMetastasis**

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

Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , obbr:response{1,3}															
Children	obbr:comment, obbr:label, obbr:response, obbr:specify															
Instance	<largestMetastasis xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </largestMetastasis>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre> &lt;xss:element name="largestMetastasis"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="size"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt; </pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

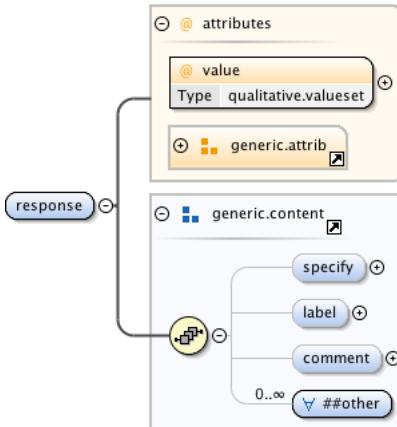
### Element nodes.status.extracapsularExtension.item / extracapsularExtension

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

Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<extracapsularExtension xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </extracapsularExtension>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>					<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use												
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>																
<b>xml:id</b>	xs:ID			optional												
Source	<pre>&lt;xss:element name="extracapsularExtension"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="nodes.extracapsularExtension.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element nodes.extracapsularExtension.response / response

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

Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<response xml:id="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </response>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:qualitative.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:qualitative.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:qualitative.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="qualitative.valueset" use="required"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element distant.section / distant

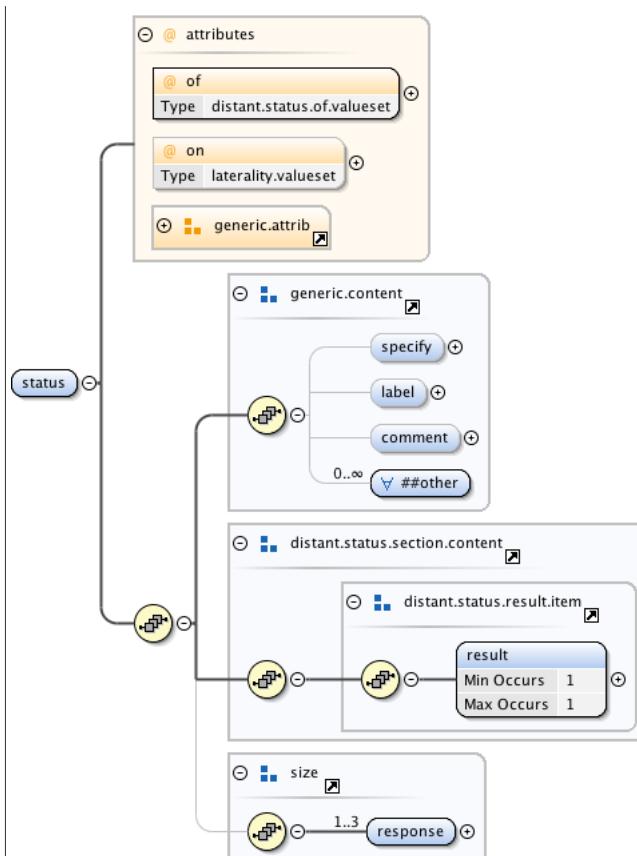
Namespace	http://purl.org/obbr/pathology/cancer/
-----------	--

Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , status															
Children	obbr:comment, obbr:label, obbr:specify, status															
Instance	<distant xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </distant>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="distant"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="distant.section.content"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

**Element distant.status.section / status**

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

## Diagram



Properties	content: complex				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , result , obbr:response{1,3}				
Children	obbr:comment, obbr:label, obbr:response, obbr:specify, result				
Instance	<status xml:id="" of="" on=""><obbr:specify>{0,1}</obbr:specify><obbr:label>{0,1}</obbr:label><obbr:comment>{0,1}</obbr:comment></status>				
Attributes	QName	Type	Fixed	Default	Use
	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>				
	<b>of</b>	obbr:distant.status.of.valueset			required
	<b>on</b>	obbr:laterality.valueset			optional
	<b>xml:id</b>	xs:ID			optional
Source	<pre> &lt;xss:element name="status"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="distant.status.section.content"/&gt;       &lt;xss:group minOccurs="0" ref="size"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attribute name="of" type="distant.status.of.valueset" use="required"/&gt;     &lt;xss:attribute name="on" type="laterality.valueset"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt; </pre>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd				

**Element `distant.status.result.item / result`**

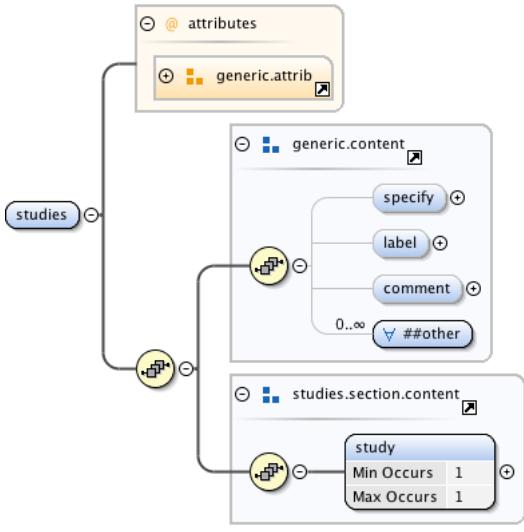
Namespace	<a href="http://purl.org/obbr/pathology/cancer/">http://purl.org/obbr/pathology/cancer/</a>																																								
Diagram	<pre> classDiagram     class result {         @ attributes         + for         + date         + generic.attrib         &lt;&gt; generic.content         &lt;&gt; distant.status.result.response     }     class generic.content {         + specify         + label         + comment         0..&gt; # other     }     class distant.status.result.response {         + response         Min Occurs 1         Max Occurs 1     }     result --&gt; generic.content     result --&gt; distant.status.result.response   </pre>																																								
Properties	content: complex																																								
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																																								
Children	obbr:comment, obbr:label, obbr:specify, response																																								
Instance	<result date="" duration="" end="" for="" xml:id="" start="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </result>																																								
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>date</b></td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>duration</b></td> <td>xs:duration</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>end</b></td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>for</b></td> <td>obbr:distant.status.result.for.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>start</b></td> <td>obbr:date.valueset</td> <td></td> <td></td> <td>optional</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					<b>date</b>	obbr:date.valueset			optional	<b>duration</b>	xs:duration			optional	<b>end</b>	obbr:date.valueset			optional	<b>for</b>	obbr:distant.status.result.for.valueset			optional	<b>start</b>	obbr:date.valueset			optional	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																																					
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																																									
<b>date</b>	obbr:date.valueset			optional																																					
<b>duration</b>	xs:duration			optional																																					
<b>end</b>	obbr:date.valueset			optional																																					
<b>for</b>	obbr:distant.status.result.for.valueset			optional																																					
<b>start</b>	obbr:date.valueset			optional																																					
<b>xml:id</b>	xs:ID			optional																																					
Source	<xs:element name="result">   <xs:complexType>     <xs:sequence>       <xs:group ref="generic.content"/>       <xs:group ref="distant.status.result.response"/>     </xs:sequence>     <xs:attribute name="for" type="distant.status.result.for.valueset"/>     <xs:attributeGroup ref="date"/>     <xs:attributeGroup ref="generic.attrib"/>   </xs:complexType> </xs:element>																																								
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																																								

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

Namespace	<a href="http://purl.org/obbr/pathology/cancer/">http://purl.org/obbr/pathology/cancer/</a>																				
Diagram	<pre> classDiagram     class response {         @ value         generic.attrib     }     class generic.content {         specify         label         comment         #other     }     response "1" -- "*" generic.content   </pre>																				
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<response xml:id="" value="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </response>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>value</b></td> <td>obbr:qualitative.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:qualitative.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:qualitative.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<xss:element name="response">   <xss:complexType>     <xss:group ref="generic.content"/>     <xss:attribute name="value" type="qualitative.valueset" use="required"/>     <xss:attributeGroup ref="generic.attrib"/>   </xss:complexType> </xss:element>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

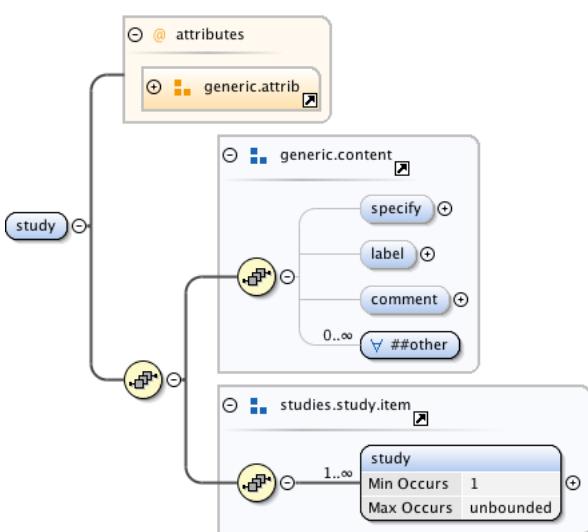
**Element `studies.section / studies`**

Namespace	<a href="http://purl.org/obbr/pathology/cancer/">http://purl.org/obbr/pathology/cancer/</a>
-----------	---

Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , study															
Children	obbr:comment, obbr:label, obbr:specify, study															
Instance	<pre>&lt;studies xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/studies&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;x:element name="studies"&gt;   &lt;x:complexType&gt;     &lt;x:sequence&gt;       &lt;x:group ref="generic.content"/&gt;       &lt;x:group ref="studies.section.content"/&gt;     &lt;/x:sequence&gt;     &lt;x:attributeGroup ref="generic.attrib"/&gt;   &lt;/x:complexType&gt; &lt;/x:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

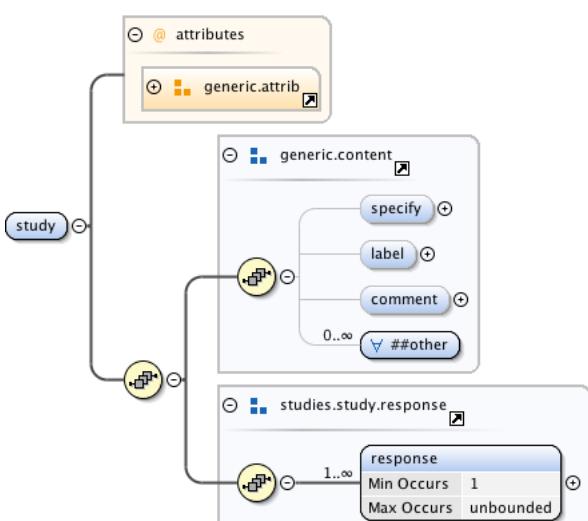
### Element studies.section.content / study

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , study+															
Children	obbr:comment, obbr:label, obbr:specify, study															
Instance	<pre>&lt;study xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/study&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="study"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="studies.study.item"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

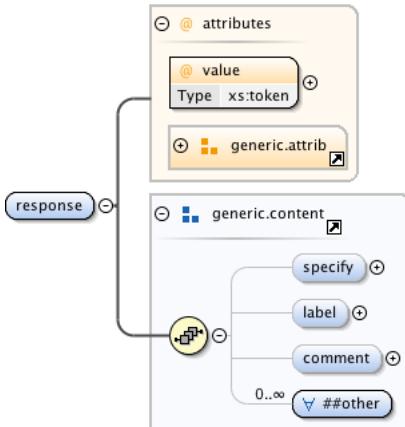
### Element studies.study.item / study

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	<p>content: complex</p> <p>maxOccurs: unbounded</p>															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer' , response+															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre>&lt;study xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/study&gt;</pre>															
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QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element maxOccurs="unbounded" name="study"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="studies.study.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element studies.study.response / response

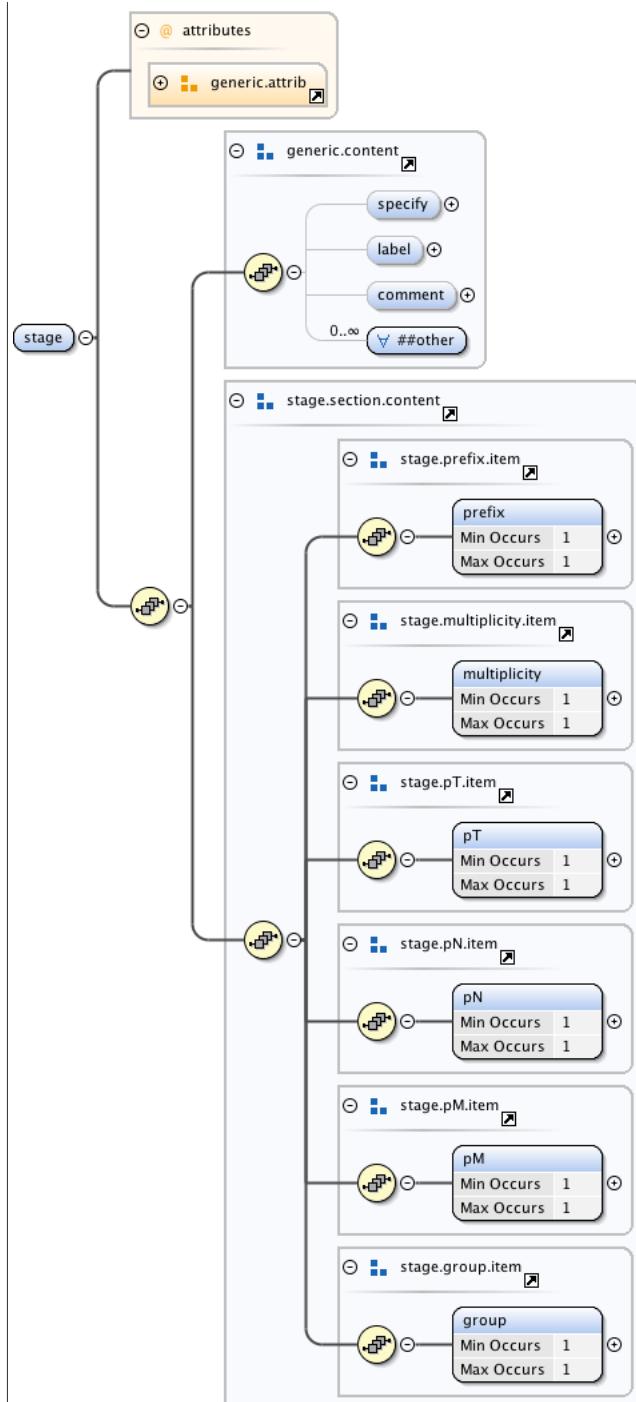
Namespace	http://purl.org/obbr/pathology/cancer/
-----------	--

Diagram																					
Properties	<p>content: complex</p> <p>maxOccurs: unbounded</p>																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
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QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	xs:token			required																	
<b>xml:id</b>	xs:ID			optional																	
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="xs:token" use="required"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

## Element stage.section / stage

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



Properties	content: complex										
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , prefix , multiplicity , pT , pN , pM , group										
Children	group, multiplicity, obbr:comment, obbr:label, obbr:specify, pM, pN, pT, prefix										
Instance	<pre> &lt;stage xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/stage&gt;   </pre>										
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/'</td><td></td><td></td><td></td><td></td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/'				
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>xml:id</b>	xs:ID			optional
Source		<xs:element name="stage"> <xs:complexType> <xs:sequence> <xs:group ref="generic.content"/> <xs:group ref="stage.section.content"/> </xs:sequence> <xs:attributeGroup ref="generic.attrib"/> </xs:complexType> </xs:element>			
Schema location		file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd			

## Element stage.prefix.item / prefix

Namespace	http://purl.org/obbr/pathology/cancer/																			
Diagram	<pre> classDiagram     class prefix {         @ attributes         generic.attrib     }     class generic.content {         specify         label         comment         0..∞ &amp;lt;--&gt; other     }     class stage.prefix.response {         response         Min Occurs: 1         Max Occurs: 1     }     prefix "1" -- "1" generic.content     prefix "1" -- "1" stage.prefix.response     generic.content "*" -- "1" other   </pre>																			
Properties	content: complex																			
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																			
Children	obbr:comment, obbr:label, obbr:specify, response																			
Instance	<prefix xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </prefix>																			
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>					QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																				
<b>xml:id</b>	xs:ID			optional																
Source	<xs:element name="prefix"> <xs:complexType> <xs:sequence> <xs:group ref="generic.content"/> <xs:group ref="stage.prefix.response"/> </xs:sequence> <xs:attributeGroup ref="generic.attrib"/> </xs:complexType> </xs:element>																			
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																			

**Element stage.prefix.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/																				
Diagram	<pre> classDiagram     class response {         @ value         generic.attrib     }     class generic.content {         specify         label         comment         #other     }     response "1" --&gt; "1" generic.content </pre>																				
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<response xml:id="" value="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </response>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>value</b></td> <td>obbr:stage.prefix.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:stage.prefix.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:stage.prefix.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<xs:element name="response">   <xs:complexType>     <xs:group ref="generic.content"/>     <xs:attribute name="value" type="stage.prefix.valueset" use="required"/>     <xs:attributeGroup ref="generic.attrib"/>   </xs:complexType> </xs:element>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

**Element stage.multiplicity.item / multiplicity**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> classDiagram     class multiplicity {         generic.attrib     }     class stage.multiplicity.response {         response     }     multiplicity "1" --&gt; "1" stage.multiplicity.response     stage.multiplicity.response "1" --&gt; "1" response     response {         Min Occurs 1         Max Occurs 1     } </pre>
Properties	content: complex
Model	response
Children	response

Instance	<multiplicity xml:id=""> <response xml:id="" value="">{1,1}</response> </multiplicity>				
Attributes	QName	Type	Fixed	Default	Use
	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>				
	<b>xml:id</b>	xs:ID			optional
Source	<xss:element name="multiplicity"> <xss:complexType> <xss:group ref="stage.multiplicity.response"/> <xss:attributeGroup ref="generic.attrib"/> </xss:complexType> </xss:element>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd				

### Element stage.multiplicity.response / response

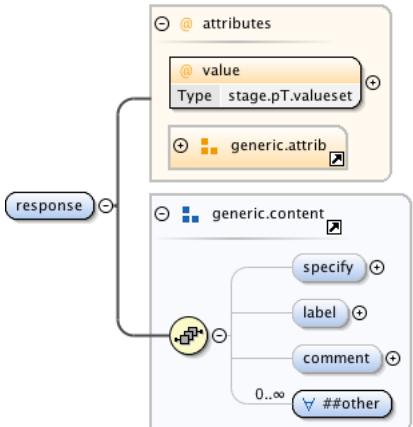
Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram	<pre> classDiagram     class response {         @ value : multiplicity.valueset         generic.attrib         &lt;&lt;generic.content&gt;&gt;             specify             label             comment             0..∞ #other     } </pre>				
Properties	content: complex				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'				
Children	obbr:comment, obbr:label, obbr:specify				
Instance	<response xml:id="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </response>				
Attributes	QName	Type	Fixed	Default	Use
	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>				
	<b>value</b>	obbr:multiplicity.valueset			required
	<b>xml:id</b>	xs:ID			optional
Source	<xss:element name="response"> <xss:complexType> <xss:group ref="generic.content"/> <xss:attribute name="value" type="multiplicity.valueset" use="required"/> <xss:attributeGroup ref="generic.attrib"/> </xss:complexType> </xss:element>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd				

**Element stage.pT.item / pT**

Namespace	http://purl.org/obbr/pathology/cancer/																			
Diagram																				
Properties	content: complex																			
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																			
Children	obbr:comment, obbr:label, obbr:specify, response																			
Instance	<pre>&lt;pT xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/pT&gt;</pre>																			
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>					QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use																
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																				
xml:id	xs:ID			optional																
Source	<pre>&lt;xsd:element name="pT"&gt;   &lt;xsd:complexType&gt;     &lt;xsd:sequence&gt;       &lt;xsd:group ref="generic.content"/&gt;       &lt;xsd:group ref="stage.pT.response"/&gt;     &lt;/xsd:sequence&gt;     &lt;xsd:attributeGroup ref="generic.attrib"/&gt;   &lt;/xsd:complexType&gt; &lt;/xsd:element&gt;</pre>																			
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																			

**Element stage.pT.response / response**

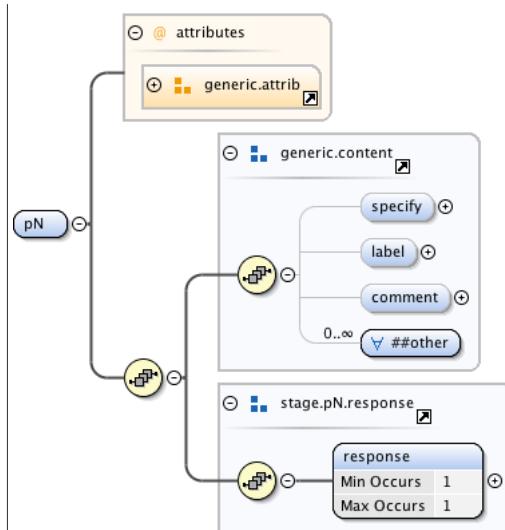
Namespace	http://purl.org/obbr/pathology/cancer/				
-----------	--	--	--	--	--

Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:stage.pT.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:stage.pT.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:stage.pT.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="stage.pT.valueset" use="required"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element stage.pN.item / pN

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



## Properties

content: complex

## Model

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

## Children

obbr:comment, obbr:label, obbr:specify, response

## Instance

```

<pN id="">
  <obbr:specify>{0,1}</obbr:specify>
  <obbr:label>{0,1}</obbr:label>
  <obbr:comment>{0,1}</obbr:comment>
</pN>

```

## Attributes

QName	Type	Fixed	Default	Use
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</b>				
<b>xml:id</b>	xs:ID			optional

## Source

```

<xss:element name="pN">
  <xss:complexType>
    <xss:sequence>
      <xss:group ref="generic.content"/>
      <xss:group ref="stage.pN.response"/>
    </xss:sequence>
    <xss:attributeGroup ref="generic.attrib"/>
  </xss:complexType>
</xss:element>

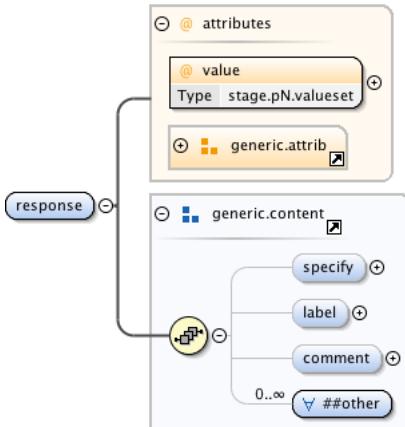
```

## Schema location

file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

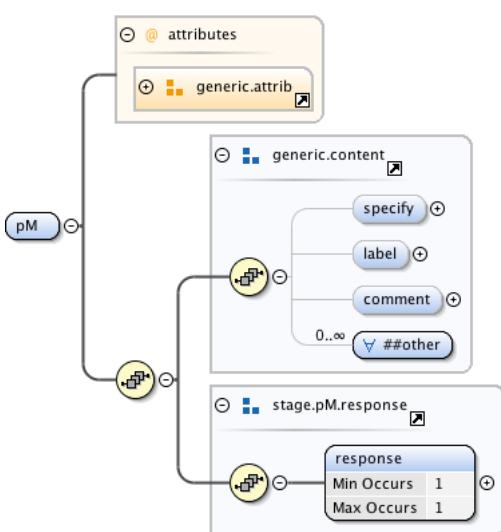
**Element stage.pN.response / response**

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:stage.pN.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:stage.pN.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:stage.pN.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xsd:element name="response"&gt;   &lt;xsd:complexType&gt;     &lt;xsd:group ref="generic.content"/&gt;     &lt;xsd:attribute name="value" type="stage.pN.valueset" use="required"/&gt;     &lt;xsd:attributeGroup ref="generic.attrib"/&gt;   &lt;/xsd:complexType&gt; &lt;/xsd:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

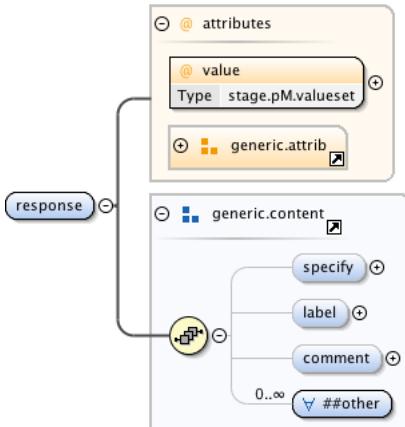
### Element stage.pM.item / pM

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pM id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </pM>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td><td></td><td></td><td></td><td></td></tr> <tr> <td>xml:id</td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre> &lt;xss:element name="pM"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="stage.pM.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt; </pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element stage.pM.response / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:stage.pM.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:stage.pM.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:stage.pM.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xsd:element name="response"&gt;   &lt;xsd:complexType&gt;     &lt;xsd:group ref="generic.content"/&gt;     &lt;xsd:attribute name="value" type="stage.pM.valueset" use="required"/&gt;     &lt;xsd:attributeGroup ref="generic.attrib"/&gt;   &lt;/xsd:complexType&gt; &lt;/xsd:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

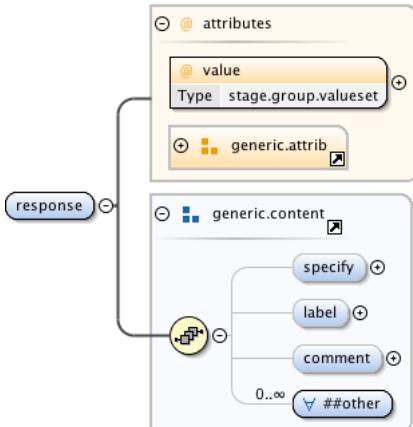
### Element stage.group.item / group

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

Diagram																
Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response															
Children	obbr:comment, obbr:label, obbr:specify, response															
Instance	<pre>&lt;group xml:id=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/group&gt;</pre>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/ obbr/pathology/ cancer/'																
xml:id	xs:ID			optional												
Source	<pre>&lt;xss:element name="group"&gt;   &lt;xss:complexType&gt;     &lt;xss:sequence&gt;       &lt;xss:group ref="generic.content"/&gt;       &lt;xss:group ref="stage.group.response"/&gt;     &lt;/xss:sequence&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

### Element stage.group.response / response

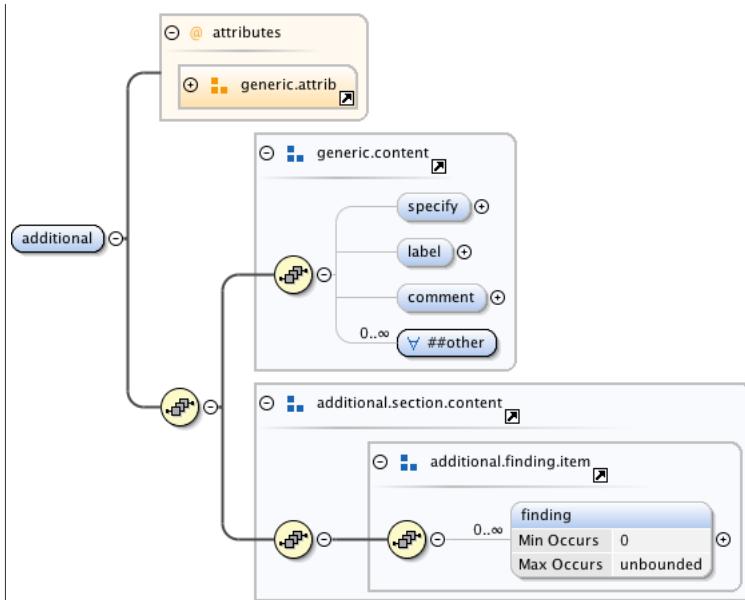
Namespace	http://purl.org/obbr/pathology/cancer/
-----------	--

Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<pre>&lt;response xml:id="" value=""&gt;   &lt;obbr:specify&gt;{0,1}&lt;/obbr:specify&gt;   &lt;obbr:label&gt;{0,1}&lt;/obbr:label&gt;   &lt;obbr:comment&gt;{0,1}&lt;/obbr:comment&gt; &lt;/response&gt;</pre>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>value</b></td><td>obbr:stage.group.valueset</td><td></td><td></td><td>required</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:stage.group.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:stage.group.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
Source	<pre>&lt;xss:element name="response"&gt;   &lt;xss:complexType&gt;     &lt;xss:group ref="generic.content"/&gt;     &lt;xss:attribute name="value" type="stage.group.valueset" use="required"/&gt;     &lt;xss:attributeGroup ref="generic.attrib"/&gt;   &lt;/xss:complexType&gt; &lt;/xss:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

### Element additional.section / additional

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



Properties	content: complex															
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , finding*															
Children	finding, obbr:comment, obbr:label, obbr:specify															
Instance	<additional xml:id=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </additional>															
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td>xml:id</td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'					xml:id	xs:ID			optional
QName	Type	Fixed	Default	Use												
ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																
xml:id	xs:ID			optional												
Source	<xss:element name="additional"> <xss:complexType> <xss:sequence> <xss:group ref="generic.content"/> <xss:group ref="additional.section.content"/> </xss:sequence> <xss:attributeGroup ref="generic.attrib"/> </xss:complexType> </xss:element>															
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd															

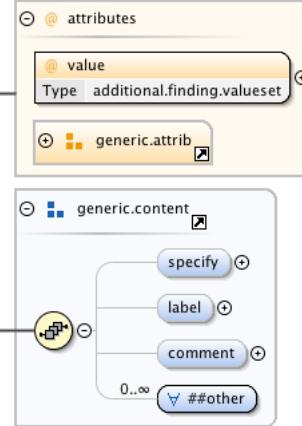
**Element additional.finding.item / finding**

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

Diagram	<pre> classDiagram     finding {         @ of Type additional.finding.of.valueset         generic.attrib     }     finding {         generic.content {             specify             label             comment         }         additional.finding.response {             response             Min Occurs 1             Max Occurs 1         }     } </pre>																				
Properties	<p>content: complex</p> <p>minOccurs: 0</p> <p>maxOccurs: unbounded</p>																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/' , response																				
Children	obbr:comment, obbr:label, obbr:specify, response																				
Instance	<finding xml:id="" of="">   <obbr:specify>{0,1}</obbr:specify>   <obbr:label>{0,1}</obbr:label>   <obbr:comment>{0,1}</obbr:comment> </finding>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th><th>Type</th><th>Fixed</th><th>Default</th><th>Use</th></tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td><td></td><td></td><td></td><td></td></tr> <tr> <td><b>of</b></td><td>obbr:additional.finding.of.valueset</td><td></td><td></td><td>optional</td></tr> <tr> <td><b>xml:id</b></td><td>xs:ID</td><td></td><td></td><td>optional</td></tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>of</b>	obbr:additional.finding.of.valueset			optional	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>of</b>	obbr:additional.finding.of.valueset			optional																	
<b>xml:id</b>	xs:ID			optional																	
Source	<xs:element maxOccurs="unbounded" minOccurs="0" name="finding">   <xs:complexType>     <xs:sequence>       <xs:group ref="generic.content"/>       <xs:group ref="additional.finding.response"/>     </xs:sequence>     <xs:attribute name="of" type="additional.finding.of.valueset"/>     <xs:attributeGroup ref="generic.attrib"/>   </xs:complexType> </xs:element>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

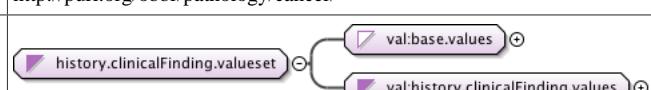
### Element additional.finding.response / response

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram																					
Properties	content: complex																				
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'																				
Children	obbr:comment, obbr:label, obbr:specify																				
Instance	<response xml:id="" value=""> <obbr:specify>{0,1}</obbr:specify> <obbr:label>{0,1}</obbr:label> <obbr:comment>{0,1}</obbr:comment> </response>																				
Attributes	<table border="1"> <thead> <tr> <th>QName</th> <th>Type</th> <th>Fixed</th> <th>Default</th> <th>Use</th> </tr> </thead> <tbody> <tr> <td><b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b></td> <td></td> <td></td> <td></td> <td></td> </tr> <tr> <td><b>value</b></td> <td>obbr:additional.finding.valueset</td> <td></td> <td></td> <td>required</td> </tr> <tr> <td><b>xml:id</b></td> <td>xs:ID</td> <td></td> <td></td> <td>optional</td> </tr> </tbody> </table>	QName	Type	Fixed	Default	Use	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>					<b>value</b>	obbr:additional.finding.valueset			required	<b>xml:id</b>	xs:ID			optional
QName	Type	Fixed	Default	Use																	
<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>																					
<b>value</b>	obbr:additional.finding.valueset			required																	
<b>xml:id</b>	xs:ID			optional																	
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="additional.finding.valueset" use="required"/&gt;     &lt;xs:attributeGroup ref="generic.attrib"/&gt;   &lt;/xs:complexType&gt; &lt;/xs:element&gt;</pre>																				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd																				

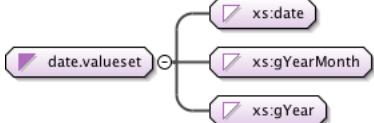
## Simple Types

### Simple Type obbr:history.clinicalFinding.valueset

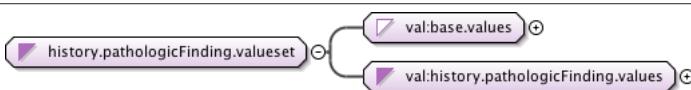
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:history.clinicalFinding.values)
Used by	Attributes history.clinicalFinding.response/response/@value, obbr:generic.content/@value
Source	<pre>&lt;xs:simpleType name="history.clinicalFinding.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:history.clinicalFinding.values"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:date.valueset

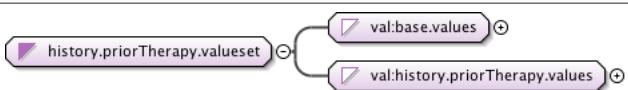
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Type	union of(xs:date, xs:gYearMonth, xs:gYear)
Used by	Attributes obbr:date/@date, obbr:date/@end, obbr:date/@start
Source	<xs:simpleType name="date.valueset"> <xs:union memberTypes="xs:date xs:gYearMonth xs:gYear" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

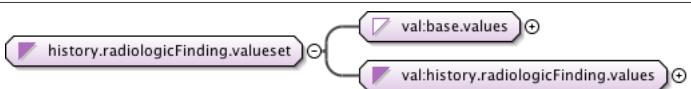
### Simple Type obbr:history.pathologicFinding.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:history.pathologicFinding.values)
Used by	Attribute history.pathologicFinding.response/response/@value
Source	<xs:simpleType name="history.pathologicFinding.valueset"> <xs:union memberTypes="val:base.values val:history.pathologicFinding.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

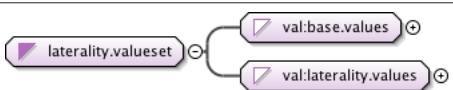
### Simple Type obbr:history.priorTherapy.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:history.priorTherapy.values)
Used by	Attribute history.priorTherapy.response/response/@value
Source	<xs:simpleType name="history.priorTherapy.valueset"> <xs:union memberTypes="val:base.values val:history.priorTherapy.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:history.radiologicFinding.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:history.radiologicFinding.values)
Used by	Attribute history.radiologicFinding.response/response/@value
Source	<xs:simpleType name="history.radiologicFinding.valueset"> <xs:union memberTypes="val:base.values val:history.radiologicFinding.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:laterality.valueset

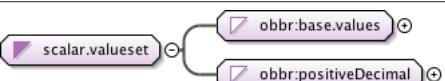
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	

Type	union of(obbr:base.values, obbr:laterality.values)	
Used by	Attributes	distant.status.section/status/@on, extent.contiguousExtension.status.result.item/result/@on, margins.margin.status.section/status/@on, nodes.status.section/status/@on, obbr:generic.content/@on, specimen.procedure.response/response/@on, tumor.site.response/response/@on
Source	<xs:simpleType name="laterality.valueset"> <xs:union memberTypes="val:base.values val:laterality.values" /> </xs:simpleType>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd	

### Simple Type obbr:specimen.procedure.response.valueset

Namespace	http://purl.org/obbr/pathology/cancer/	
Diagram		
Type	union of(obbr:base.values)	
Used by	Attribute specimen.procedure.response/response/@value	
Source	<xs:simpleType name="specimen.procedure.response.valueset"> <xs:union memberTypes="val:base.values" /> </xs:simpleType>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd	

### Simple Type obbr:scalar.valueset

Namespace	http://purl.org/obbr/pathology/cancer/	
Diagram		
Type	union of(obbr:base.values, obbr:positiveDecimal)	
Used by	Attributes obbr:quantity/@value, obbr:weight/@value	
Source	<xs:simpleType name="scalar.valueset"> <xs:union memberTypes="obbr:base.values obbr:positiveDecimal" /> </xs:simpleType>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd	

### Simple Type obbr:lengthUnit.valueset

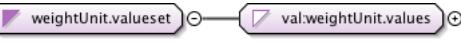
Namespace	http://purl.org/obbr/pathology/cancer/	
Diagram		
Type	union of(obbr:lengthUnit.values)	
Used by	Attribute obbr:quantity/@unit	
Source	<xs:simpleType name="lengthUnit.valueset"> <xs:union memberTypes="val:lengthUnit.values" /> </xs:simpleType>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd	

### Simple Type obbr:relation.valueset

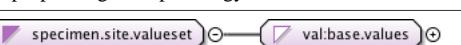
Namespace	http://purl.org/obbr/pathology/cancer/	
Diagram		
Type	union of(obbr:relation.values)	
Used by	Attributes obbr:quantity/@relation, obbr:tally/@relation, obbr:weight/@relation	
Source	<xs:simpleType name="relation.valueset"> <xs:union memberTypes="val:relation.values" /> </xs:simpleType>	

Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd
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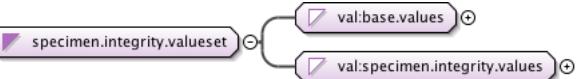
### Simple Type obbr:weightUnit.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:weightUnit.values)
Used by	Attribute obbr:weight/@unit
Source	<xs:simpleType name="weightUnit.valueset"> <xs:union memberTypes="val:weightUnit.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

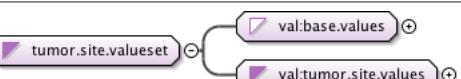
### Simple Type obbr:specimen.site.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute specimen.site.response/response/@value
Source	<xs:simpleType name="specimen.site.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:specimen.integrity.valueset

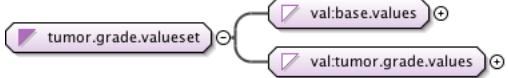
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:specimen.integrity.values)
Used by	Attribute specimen.integrity.response/response/@value
Source	<xs:simpleType name="specimen.integrity.valueset"> <xs:union memberTypes="val:base.values val:specimen.integrity.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:tumor.site.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:tumor.site.values)
Used by	Attribute tumor.site.response/response/@value
Source	<xs:simpleType name="tumor.site.valueset"> <xs:union memberTypes="val:base.values val:tumor.site.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:tumor.grade.valueset

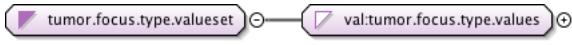
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Type	union of(obbr:base.values, obbr:tumor.grade.values)
Used by	Attribute tumor.grade.response/response/@value
Source	<xs:simpleType name="tumor.grade.valueset"> <xs:union memberTypes="val:base.values val:tumor.grade.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:tumor.gradeSystem.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:tumor.gradeSystem.values)
Used by	Attribute tumor.grade.item/grade/@system
Source	<xs:simpleType name="tumor.gradeSystem.valueset"> <xs:union memberTypes="val:base.values val:tumor.gradeSystem.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:tumor.focus.type.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:tumor.focus.type.values)
Used by	Attribute tumor.focus.section/focus/@type
Source	<xs:simpleType name="tumor.focus.type.valueset"> <xs:union memberTypes="val:tumor.focus.type.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:focality.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:focality.values)
Used by	Attributes extent.contiguousExtension.status.result.response/response/@extent, margins.margin.status.result.response/response/@extent, obbr:generic.content/@extent
Source	<xs:simpleType name="focality.valueset"> <xs:union memberTypes="val:focality.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Simple Type obbr:qualitative.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:qualitative.values)

Used by	Attributes	distant.status.result.response/response/@value, extent.contiguousExtension.status.result.response/response/ @value, margins.margin.status.result.response/response/ @closest, margins.margin.status.result.response/response/@value, nodes.extracapsularExtension.response/response/@value, obbr:generic.content/@closest
Source		<xs:simpleType name="qualitative.valueset"> <xs:union memberTypes="val:base.values val:qualitative.values" /> </xs:simpleType>
Schema location		file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

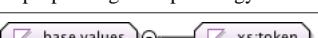
### Simple Type obbr:extent.contiguousExtension.status.result.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:extent.contiguousExtension.status.result.for.values)
Used by	Attribute extent.contiguousExtension.status.result.item/result/@for
Source	<xs:simpleType name="extent.contiguousExtension.status.result.for.valueset"> <xs:union memberTypes="val:extent.contiguousExtension.status.result.for.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

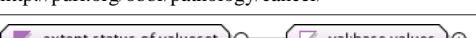
### Simple Type obbr:extent.contiguousExtension.status.distance.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:extent.contiguousExtension.status.distance.for.values)
Used by	Attribute extent.contiguousExtension.status.distance.item/distance/@for
Source	<xs:simpleType name="extent.contiguousExtension.status.distance.for.valueset"> <xs:union memberTypes="val:extent.contiguousExtension.status.distance.for.values" /> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:base.values

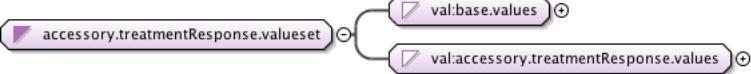
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration specify enumeration inapplicable enumeration unreported
Used by	Attribute extent.contiguousExtension.status.distance.item/distance/@from
Source	<xs:simpleType name="base.values"> <xs:restriction base="xs:token"> <xs:enumeration value="specify"/> <xs:enumeration value="inapplicable"/> <xs:enumeration value="unreported"/> </xs:restriction> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

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

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

Type	union of(obbr:base.values)
Used by	Attribute extent.contiguousExtension.status.section/status/@of
Source	<xs:simpleType name="extent.status.of.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:accessory.treatmentResponse.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:accessory.treatmentResponse.values)
Used by	Attribute accessory.treatmentResponse.response/response/@value
Source	<xs:simpleType name="accessory.treatmentResponse.valueset"> <xs:union memberTypes="val:base.values val:accessory.treatmentResponse.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:accessory.lymphvascularInvasion.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:qualitative.values)
Used by	Attribute accessory.lymphvascularInvasion.response/response/@value
Source	<xs:simpleType name="accessory.lymphvascularInvasion.valueset"> <xs:union memberTypes="val:base.values val:qualitative.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:accessory.perineuralInvasion.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:qualitative.values)
Used by	Attribute accessory.perineuralInvasion.response/response/@value
Source	<xs:simpleType name="accessory.perineuralInvasion.valueset"> <xs:union memberTypes="val:base.values val:qualitative.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:margins.margin.status.result.for.valueset

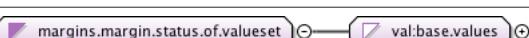
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:margins.margin.status.result.for.values)
Used by	Attribute margins.margin.status.result.item/result/@for

Source	<pre>&lt;xs:simpleType name="margins.margin.status.result.for.valueset"&gt;   &lt;xs:union memberTypes="val:base.values val:margins.margin.status.result.for.values"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

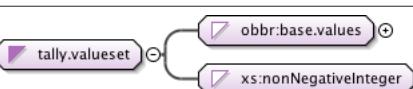
### Simple Type obbr:margins.margin.distance.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:margins.margin.status.result.for.valueset)
Used by	Attribute margins.margin.distance.item/distance/@for
Source	<pre>&lt;xs:simpleType name="margins.margin.distance.for.valueset"&gt;   &lt;xs:union memberTypes="val:margins.margin.status.result.for.valueset"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:margins.margin.status.of.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute margins.margin.status.section/status/@of
Source	<pre>&lt;xs:simpleType name="margins.margin.status.of.valueset"&gt;   &lt;xs:union memberTypes="val:base.values"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:tally.valueset

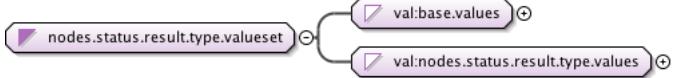
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, xs:nonNegativeInteger)
Used by	Attribute obbr:tally/@value
Source	<pre>&lt;xs:simpleType name="tally.valueset"&gt;   &lt;xs:union memberTypes="obbr:base.values xs:nonNegativeInteger"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

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

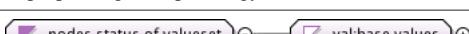
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:nodes.status.result.for.values)
Used by	Attribute nodes.status.result.section/result/@for
Source	<pre>&lt;xs:simpleType name="nodes.status.result.for.valueset"&gt;   &lt;xs:union memberTypes="val:nodes.status.result.for.values"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

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

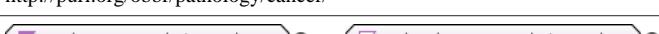
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Type	union of(obbr:base.values, obbr:nodes.status.result.type.values)
Used by	Attribute nodes.status.result.section/result/@type
Source	<xs:simpleType name="nodes.status.result.type.valueset"> <xs:union memberTypes="val:base.values val:nodes.status.result.type.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

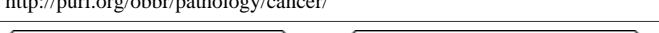
### Simple Type obbr:status.of.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute nodes.status.section/status/@of
Source	<xs:simpleType name="nodes.status.of.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

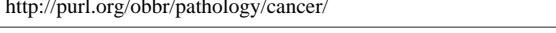
### Simple Type obbr:status.technique.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:status.technique.values)
Used by	Attribute nodes.status.section/status/@technique
Source	<xs:simpleType name="nodes.status.technique.valueset"> <xs:union memberTypes="val:nodes.status.technique.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

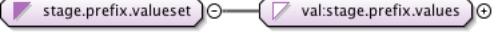
### Simple Type obbr:distant.status.result.for.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:distant.status.result.for.values)
Used by	Attribute distant.status.result.item/result/@for
Source	<xs:simpleType name="distant.status.result.for.valueset"> <xs:union memberTypes="val:distant.status.result.for.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

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

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:distant.status.of.values)
Used by	Attribute distant.status.section/status/@of
Source	<xs:simpleType name="distant.status.of.valueset"> <xs:union memberTypes="val:distant.status.of.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:stage.prefix.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:stage.prefix.values)
Used by	Attribute stage.prefix.response/response/@value
Source	<xs:simpleType name="stage.prefix.valueset"> <xs:union memberTypes="val:stage.prefix.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:multiplicity.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:multiplicity.values)
Used by	Attribute stage.multiplicity.response/response/@value
Source	<xs:simpleType name="multiplicity.valueset"> <xs:union memberTypes="val:multiplicity.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Simple Type obbr:stage.pT.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute stage.pT.response/response/@value
Source	<xs:simpleType name="stage.pT.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:stage.pN.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute stage.pN.response/response/@value
Source	<xs:simpleType name="stage.pN.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:stage.pM.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute stage.pM.response/response/@value
Source	<xs:simpleType name="stage.pM.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>

	</xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:stage.group.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute stage.group.response/response/@value
Source	<xs:simpleType name="stage.group.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

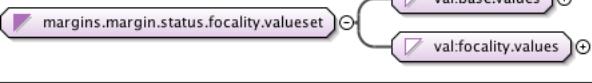
### Simple Type obbr:additional.finding.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute additional.finding.response/response/@value
Source	<xs:simpleType name="additional.finding.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:additional.finding.of.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Used by	Attribute additional.finding.item/finding/@of
Source	<xs:simpleType name="additional.finding.of.valueset"> <xs:union memberTypes="val:base.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:margins.margin.status.focality.valueset

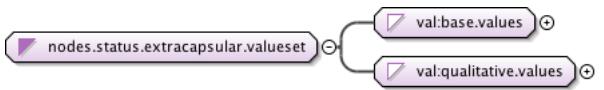
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:focality.values)
Source	<xs:simpleType name="margins.margin.status.focality.valueset"> <xs:union memberTypes="val:base.values val:focality.values"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:margins.margin.distance.from.valueset

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

Source	<xss:simpleType name="margins.margin.distance.from.valueset"> <xss:union memberTypes="val:base.values"/> </xss:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

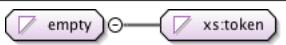
### Simple Type obbr:nodes.status.extracapsular.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values, obbr:qualitative.values)
Source	<xss:simpleType name="nodes.status.extracapsular.valueset"> <xss:union memberTypes="val:base.values val:qualitative.values"/> </xss:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

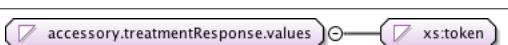
### Simple Type obbr:tumor.histologicType.valueset

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:base.values)
Source	<xss:simpleType name="tumor.histologicType.valueset"> <xss:union memberTypes="val:base.values"/> </xss:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:empty

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration unreported
Source	<xss:simpleType name="empty"> <xss:restriction base="xs:token"> <xss:enumeration value="unreported"/> </xss:restriction> </xss:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/valuesets.xsd

### Simple Type obbr:accessory.treatmentResponse.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration none enumeration slight enumeration marked enumeration complete
Source	<xss:simpleType name="accessory.treatmentResponse.values"> <xss:restriction base="xs:token"> <xss:enumeration value="none"/> <xss:enumeration value="slight"/> <xss:enumeration value="marked"/> <xss:enumeration value="complete"/> </xss:restriction> </xss:simpleType>

Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd
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### Simple Type obbr:extent.contiguousExtension.status.result.for.values

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Type	restriction of xs:token				
Facets	<table> <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>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd				

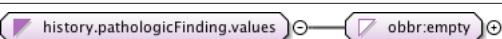
### Simple Type obbr:extent.contiguousExtension.status.distance.for.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:extent.contiguousExtension.status.result.for.values)
Source	<pre>&lt;xs:simpleType name="extent.contiguousExtension.status.distance.for.values"&gt;   &lt;xs:union memberTypes="obbr:extent.contiguousExtension.status.result.for.values"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

### Simple Type obbr:history.priorTherapy.values

Namespace	http://purl.org/obbr/pathology/cancer/						
Diagram							
Type	restriction of xs:token						
Facets	<table> <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>						
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd						

### Simple Type obbr:history.pathologicFinding.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:empty)
Source	<pre>&lt;xs:simpleType name="history.pathologicFinding.values"&gt;   &lt;xs:union memberTypes="obbr:empty"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

### Simple Type obbr:history.clinicalFinding.values

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Type	union of(obbr:empty)
Source	<xs:simpleType name="history.clinicalFinding.values"> <xs:union memberTypes="obbr:empty"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

### Simple Type obbr:history.radiologicFinding.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:empty)
Source	<xs:simpleType name="history.radiologicFinding.values"> <xs:union memberTypes="obbr:empty"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

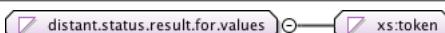
### Simple Type obbr:margins.margin.status.result.for.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration      in-situ enumeration      invasive
Source	<xs:simpleType name="margins.margin.status.result.for.values"> <xs:restriction base="xs:token"> <xs:enumeration value="in-situ"/> <xs:enumeration value="invasive"/> </xs:restriction> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

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

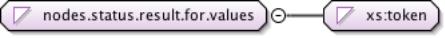
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration      liver enumeration      bone enumeration      lung enumeration      brain
Source	<xs:simpleType name="distant.status.of.values"> <xs:restriction base="xs:token"> <xs:enumeration value="liver"/> <xs:enumeration value="bone"/> <xs:enumeration value="lung"/> <xs:enumeration value="brain"/> </xs:restriction> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

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

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

Type	restriction of xs:token
Facets	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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

### Simple Type obbr:node.status.result.for.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

### Simple Type obbr:node.status.result.type.values

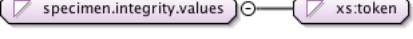
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration necroinflammatory
Source	<pre>&lt;xs:simpleType name="nodes.status.result.type.values"&gt;   &lt;xs:restriction base="xs:token"&gt;     &lt;xs:enumeration value="necroinflammatory"/&gt;   &lt;/xs:restriction&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

### Simple Type obbr:node.status.technique.values

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

	</xs:restriction> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

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

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Type	restriction of xs:token				
Facets	<table> <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>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd				

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

Namespace	http://purl.org/obbr/pathology/cancer/								
Diagram									
Type	restriction of xs:token								
Facets	<table> <tr> <td>enumeration</td> <td>y</td> </tr> <tr> <td>enumeration</td> <td>r</td> </tr> <tr> <td>enumeration</td> <td>a</td> </tr> <tr> <td>enumeration</td> <td>none</td> </tr> </table>	enumeration	y	enumeration	r	enumeration	a	enumeration	none
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>								
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd								

### Simple Type obbr:stage.pT.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:empty)
Source	<pre>&lt;xs:simpleType name="stage.pT.values"&gt;   &lt;xs:union memberTypes="obbr:empty"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

### Simple Type obbr:stage.pN.values

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

Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd
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### Simple Type obbr:stage.pM.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:empty)
Source	<xs:simpleType name="stage.pM.values"> <xs:union memberTypes="obbr:empty"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

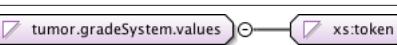
### Simple Type obbr:stage.group.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:empty)
Source	<xs:simpleType name="stage.group.values"> <xs:union memberTypes="obbr:empty"/> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

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

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration      1 enumeration      2 enumeration      3 enumeration      4 enumeration      X
Source	<xs:simpleType name="tumor.grade.values"> <xs:restriction base="xs:token"> <xs:enumeration value="1"/> <xs:enumeration value="2"/> <xs:enumeration value="3"/> <xs:enumeration value="4"/> <xs:enumeration value="X"/> </xs:restriction> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

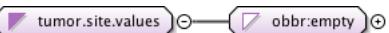
### Simple Type obbr:tumor.gradeSystem.values

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:token
Facets	enumeration      4-grade
Source	<xs:simpleType name="tumor.gradeSystem.values"> <xs:restriction base="xs:token"> <xs:enumeration value="4-grade"/> </xs:restriction> </xs:simpleType>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

**Simple Type obbr:tumor.focus.type.values**

Namespace	http://purl.org/obbr/pathology/cancer/						
Diagram							
Type	restriction of xs:token						
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						
Source	<pre>&lt;xs:simpleType name="tumor.focus.type.values"&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;</pre>						
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd						

**Simple Type obbr:tumor.site.values**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	union of(obbr:empty)
Source	<pre>&lt;xs:simpleType name="tumor.site.values"&gt;   &lt;xs:union memberTypes="obbr:empty"/&gt; &lt;/xs:simpleType&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/values.xsd

**Simple Type obbr:positiveDecimal**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Type	restriction of xs:decimal
Facets	minInclusive 0.0
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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Simple Type obbr:focality.values**

Namespace	http://purl.org/obbr/pathology/cancer/								
Diagram									
Type	restriction of xs:token								
Facets	<table> <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>								
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd								

### Simple Type obbr:laterality.values

Namespace	http://purl.org/obbr/pathology/cancer/												
Diagram													
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>												
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd												

### Simple Type obbr:lengthUnit.values

Namespace	http://purl.org/obbr/pathology/cancer/						
Diagram							
Type	restriction of xs:token						
Facets	<table> <tr><td>enumeration</td><td>cm</td></tr> <tr><td>enumeration</td><td>mm</td></tr> <tr><td>enumeration</td><td>in</td></tr> </table>	enumeration	cm	enumeration	mm	enumeration	in
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>						
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd						

### Simple Type obbr:multiplicity.values

Namespace	http://purl.org/obbr/pathology/cancer/										
Diagram											
Type	restriction of xs:token										
Facets	<table> <tr><td>enumeration</td><td>1</td></tr> <tr><td>enumeration</td><td>2</td></tr> <tr><td>enumeration</td><td>3</td></tr> <tr><td>enumeration</td><td>4</td></tr> <tr><td>enumeration</td><td>m</td></tr> </table>	enumeration	1	enumeration	2	enumeration	3	enumeration	4	enumeration	m
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>										

Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd
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### Simple Type obbr:qualitative.values

Namespace	http://purl.org/obbr/pathology/cancer/								
Diagram									
Type	restriction of xs:token								
Facets	<table> <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	positive	enumeration	negative	enumeration	equivocal	enumeration	indeterminate
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>								
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd								

### Simple Type obbr:relation.values

Namespace	http://purl.org/obbr/pathology/cancer/												
Diagram													
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>												
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd												

### Simple Type obbr:weightUnit.values

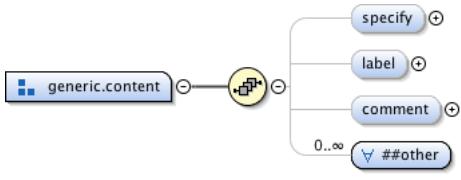
Namespace	http://purl.org/obbr/pathology/cancer/						
Diagram							
Type	restriction of xs:token						
Facets	<table> <tr> <td>enumeration</td> <td>mg</td> </tr> <tr> <td>enumeration</td> <td>g</td> </tr> <tr> <td>enumeration</td> <td>kg</td> </tr> </table>	enumeration	mg	enumeration	g	enumeration	kg
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>						

Schema location

file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

## Element Groups

### Element Group obbr:generic.content

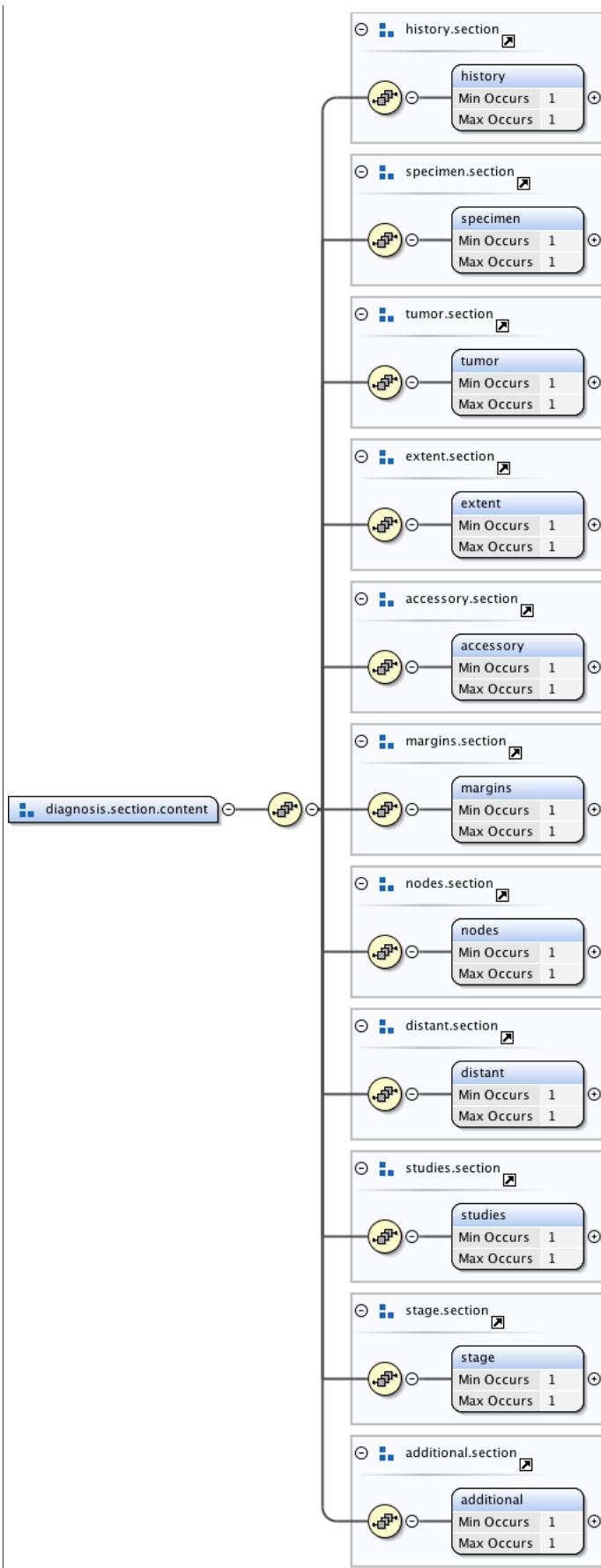
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	 <pre> classDiagram     class generic.content     class other {         &lt;&lt;#other&gt;&gt;     }     generic.content "0..1" -- "0..1" specify     generic.content "0..1" -- "0..1" label     generic.content "0..1" -- "0..infinity" #other   </pre>
Used by	<p>Elements</p> <p>accessory.lymphvascularInvasion.item/lymphvascularInvasion,  accessory.lymphvascularInvasion.response/response, accessory.perineuralInvasion.item/  perineuralInvasion, accessory.perineuralInvasion.response/response,  accessory.section/accessory, accessory.treatmentResponse.item/treatmentResponse,  accessory.treatmentResponse.response/response, additional.finding.item/finding,  additional.finding.response/response, additional.section/additional, diagnosis,  distant.section/distant, distant.status.result.item/result, distant.status.result.response/  response, distant.status.section/status, extent.contiguousExtension.section/  contiguousExtension, extent.contiguousExtension.status.distance.item/  distance, extent.contiguousExtension.status.result.item/result,  extent.contiguousExtension.status.result.response/response,  extent.contiguousExtension.status.section/status, extent.section/extent,  history.clinicalFinding.item/clinicalFinding, history.clinicalFinding.response/response,  history.pathologicFinding.item/pathologicFinding, history.pathologicFinding.response/  response, history.priorTherapy.item/priorTherapy, history.priorTherapy.response/response,  history.radiologicFinding.item/radiologicFinding, history.radiologicFinding.response/  response, history.section/history, margins.margin.distance.item/distance,  margins.margin.section/margin, margins.margin.status.result.item/result,  margins.margin.status.result.response/response, margins.margin.status.section/  status, margins.section/margins, nodes.extracapsularExtension.response/response,  nodes.section/nodes, nodes.status.extracapsularExtension.item/extracapsularExtension,  nodes.status.largestMetastasis.item/largestMetastasis, nodes.status.result.response/response,  nodes.status.result.section/result, nodes.status.section/status, obbr:length/obbr:response,  obbr:size/obbr:response, specimen.integrity.item/integrity, specimen.integrity.response/  response, specimen.procedure.item/procedure, specimen.procedure.response/response,  specimen.section/specimen, specimen.site.item/site, specimen.site.response/response,  specimen.size.item/size, specimen.weight.item/weight, specimen.weight.response/response,  stage.group.item/group, stage.group.response/response, stage.multiplicity.response/response,  stage.pM.item/pM, stage.pM.response/response, stage.pN.item/pN, stage.pN.response/  response, stage.pT.item/pT, stage.pT.response/response, stage.prefix.item/prefix,  stage.prefix.response/response, stage.section/stage, studies.section.content/study,  studies.section/studies, studies.study.item/study, studies.study.response/response,  tumor.focus.section/focus, tumor.grade.item/grade, tumor.grade.response/response,  tumor.histologicType.item/histologicType, tumor.histologicType.response/response,  tumor.section/tumor, tumor.site.item/site, tumor.site.response/response, tumor.size.item/size</p>
Model	obbr:specify{0,1} , obbr:label{0,1} , obbr:comment{0,1} , ANY element from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'
Children	obbr:comment, obbr:label, obbr:specify
Source	<pre> &lt;xss:group name="generic.content"&gt;   &lt;xss:sequence&gt;     &lt;xss:element minOccurs="0" name="specify" type="xs:string"/&gt;     &lt;xss:element minOccurs="0" name="label" type="xs:string"/&gt;     &lt;xss:element minOccurs="0" name="comment" type="xs:string"/&gt;     &lt;xss:any maxOccurs="unbounded" minOccurs="0" namespace="#other"       processContents="skip"/&gt;   &lt;/xss:sequence&gt; &lt;/xss:group&gt;   </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Element Group diagnosis.section.content

Namespace

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

## Diagram



Used by

Element diagnosis

Model	history , specimen , tumor , extent , accessory , margins , nodes , distant , studies , stage , additional
Children	accessory, additional, distant, extent, history, margins, nodes, specimen, stage, studies, tumor
Source	<pre>&lt;xs:group name="diagnosis.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="history.section"/&gt;     &lt;xs:group ref="specimen.section"/&gt;     &lt;xs:group ref="tumor.section"/&gt;     &lt;xs:group ref="extent.section"/&gt;     &lt;xs:group ref="accessory.section"/&gt;     &lt;xs:group ref="margins.section"/&gt;     &lt;xs:group ref="nodes.section"/&gt;     &lt;xs:group ref="distant.section"/&gt;     &lt;xs:group ref="studies.section"/&gt;     &lt;xs:group ref="stage.section"/&gt;     &lt;xs:group ref="additional.section"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

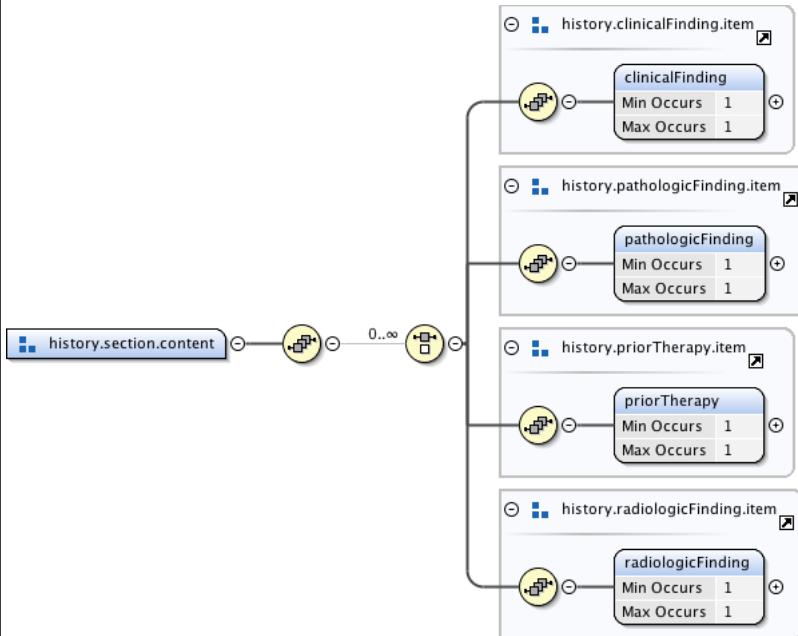
### Element Group **history.section**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> classDiagram     class historySection {         &lt;&lt;history&gt;&gt;         &lt;&lt;Min Occurs 1&gt;&gt;         &lt;&lt;Max Occurs 1&gt;&gt;     }     historySection &lt; -- history     historySection &lt; -- genericContent     historySection &lt; -- historySectionContent     historySection &lt; -- genericAttrib   </pre>
Used by	Element Group diagnosis.section.content
Model	history
Children	history
Source	<pre>&lt;xs:group name="history.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="history"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="history.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group **history.section.content**

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



## Used by

Element history.section/history

## Model

((clinicalFinding) | (pathologicFinding) | (priorTherapy) | (radiologicFinding))

## Children

clinicalFinding, pathologicFinding, priorTherapy, radiologicFinding

## Source

```

<xs:group name="history.section.content">
  <xs:sequence>
    <xs:choice maxOccurs="unbounded" minOccurs="0">
      <xs:group ref="history.clinicalFinding.item"/>
      <xs:group ref="history.pathologicFinding.item"/>
      <xs:group ref="history.priorTherapy.item"/>
      <xs:group ref="history.radiologicFinding.item"/>
    </xs:choice>
  </xs:sequence>
</xs:group>
  
```

## Schema location

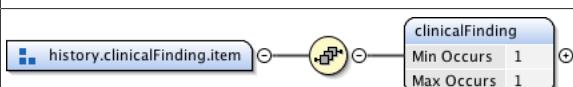
file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element Group history.clinicalFinding.item**

## Namespace

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

## Diagram



## Used by

Element Group history.section.content

## Model

clinicalFinding

## Children

clinicalFinding

## Source

```

<xs:group name="history.clinicalFinding.item">
  <xs:sequence>
    <xs:element name="clinicalFinding">
      <xs:complexType>
        <xs:sequence>
          <xs:group ref="generic.content"/>
          <xs:group ref="history.clinicalFinding.response"/>
        </xs:sequence>
        <xs:attributeGroup ref="date"/>
        <xs:attributeGroup ref="generic.attrib"/>
      </xs:complexType>
    </xs:element>
  </xs:sequence>
</xs:group>
  
```

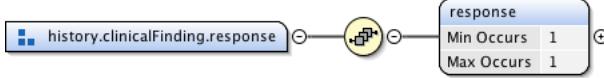
## Schema location

file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

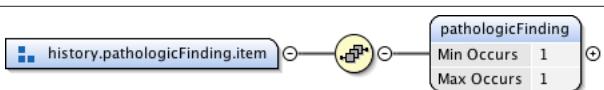
**Element Group history.clinicalFinding.response**

## Namespace

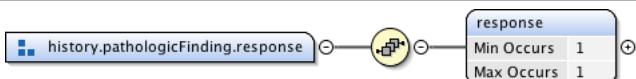
http://purl.org/obbr/pathology/cancer/

Diagram	
Used by	Element history.clinicalFinding.item/clinicalFinding
Model	response
Children	response
Source	<pre>&lt;xs:group name="history.clinicalFinding.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="history.clinicalFinding.valueset" use="required"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group history.pathologicFinding.item

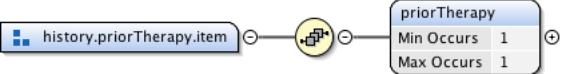
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group history.section.content
Model	pathologicFinding
Children	pathologicFinding
Source	<pre>&lt;xs:group name="history.pathologicFinding.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="pathologicFinding"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="history.pathologicFinding.response"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="date"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group history.pathologicFinding.response

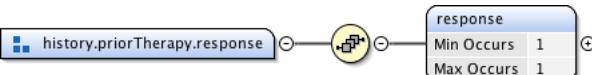
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element history.pathologicFinding.item/pathologicFinding
Model	response
Children	response
Source	<pre>&lt;xs:group name="history.pathologicFinding.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;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

	</xs:sequence> </xs:group>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

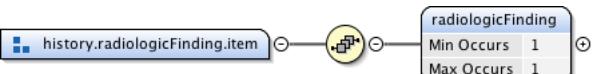
### Element Group history.priorTherapy.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group history.section.content
Model	priorTherapy
Children	priorTherapy
Source	<pre>&lt;xs:group name="history.priorTherapy.item"&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="history.priorTherapy.response"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attributeGroup ref="date"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group history.priorTherapy.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element history.priorTherapy.item/priorTherapy
Model	response
Children	response
Source	<pre>&lt;xs:group name="history.priorTherapy.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="history.priorTherapy.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group history.radiologicFinding.item

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

	<pre> &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="history.radiologicFinding.response"/&gt;       &lt;/xs:sequence&gt;       &lt;xs:attributeGroup ref="date"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group history.radiologicFinding.response

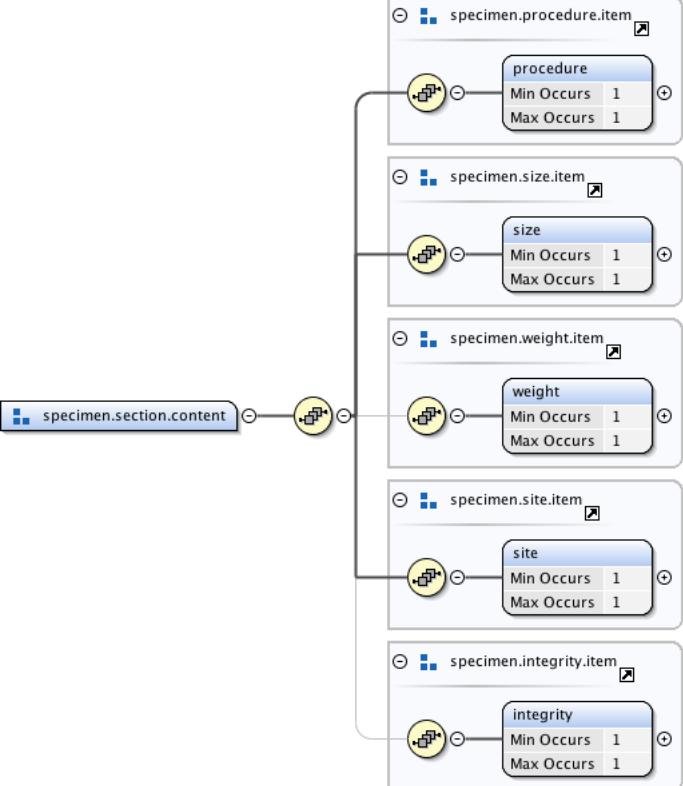
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[history.radiologicFinding.response] --&gt; B(( ))     B --&gt; C[response]     C -- Min Occurs 1 --&gt; D[ ]     C -- Max Occurs 1 --&gt; E[ ] </pre>
Used by	Element history.radiologicFinding.item/radiologicFinding
Model	response
Children	response
Source	<pre> &lt;xs:group name="history.radiologicFinding.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="history.radiologicFinding.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.section

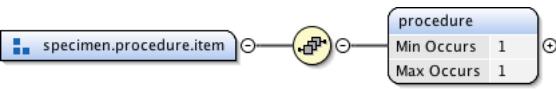
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[specimen.section] --&gt; B(( ))     B --&gt; C[specimen]     C -- Min Occurs 1 --&gt; D[ ]     C -- Max Occurs 1 --&gt; E[ ] </pre>
Used by	Element Group diagnosis.section.content
Model	specimen
Children	specimen
Source	<pre> &lt;xs:group name="specimen.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="specimen"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="specimen.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.section.content

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element specimen.section/specimen
Model	procedure , size , weight , site , integrity
Children	integrity, procedure, site, size, weight
Source	<pre>&lt;xs:group name="specimen.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="specimen.procedure.item"/&gt;     &lt;xs:group ref="specimen.size.item"/&gt;     &lt;xs:group minOccurs="0" ref="specimen.weight.item"/&gt;     &lt;xs:group ref="specimen.site.item"/&gt;     &lt;xs:group minOccurs="0" ref="specimen.integrity.item"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.procedure.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group specimen.section.content
Model	procedure
Children	procedure
Source	<pre>&lt;xs:group name="specimen.procedure.item"&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.response"/&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>

Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd
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### Element Group specimen.procedure.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[specimen.procedure.response] --- B(( ))     B --- C[response]     C --- D[Min Occurs: 1]     C --- E[Max Occurs: unbounded]   </pre>
Used by	Element specimen.procedure.item/procedure
Model	response+
Children	response
Source	<pre> &lt;xs:group name="specimen.procedure.response"&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:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;   </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.size.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[specimen.size.item] --- B(( ))     B --- C[size]     C --- D[Min Occurs: 1]     C --- E[Max Occurs: 1]   </pre>
Used by	Element Group specimen.section.content
Model	size
Children	size
Source	<pre> &lt;xs:group name="specimen.size.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="size"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="specimen.size.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.size.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[specimen.size.response] --- B(( ))     B --- C[size]     C --- D[specimen.size.item]     C --- E[response]     B --- F(( ))     F --- G[1..3]     F --- H[response]   </pre>
Used by	Element specimen.size.item/size
Model	obbr:response{1,3}
Children	obbr:response
Source	<pre> &lt;xs:group name="specimen.size.response"&gt;   &lt;xs:sequence&gt;   </pre>

	<pre> &lt;xs:group ref="size" /&gt; &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group obbr:size

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     size["size"] -- "1..3" --&gt; response["response"] </pre>
Used by	Element Groups specimen.size.response, tumor.size.response Elements distant.status.section/status, nodes.status.largestMetastasis.item/largestMetastasis
Model	obbr:response{1,3}
Children	obbr: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="obbr:generic.content"/&gt;         &lt;xs:attributeGroup ref="obbr:quantity"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Element Group specimen.weight.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     item["specimen.weight.item"] -- "1..3" --&gt; weight["weight Min Occurs 1 Max Occurs 1"] </pre>
Used by	Element Group specimen.section.content
Model	weight
Children	weight
Source	<pre> &lt;xs:group name="specimen.weight.item"&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.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.weight.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     item["specimen.weight.response"] -- "1..3" --&gt; response["response Min Occurs 1 Max Occurs 1"] </pre>
Used by	Element specimen.weight.item/weight
Model	response
Children	response
Source	<pre> &lt;xs:group name="specimen.weight.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt; </pre>

	<pre> &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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.site.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[specimen.site.item] --&gt; B[site]     B -- "Min Occurs 1, Max Occurs 1" --&gt; C </pre>
Used by	Element Group specimen.section.content
Model	site
Children	site
Source	<pre> &lt;xs:group name="specimen.site.item"&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="specimen.site.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.site.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[specimen.site.response] --&gt; B[response]     B -- "Min Occurs 1, Max Occurs unbounded" --&gt; C </pre>
Used by	Element specimen.site.item/site
Model	response+
Children	response
Source	<pre> &lt;xs:group name="specimen.site.response"&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.site.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.integrity.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[specimen.integrity.item] --&gt; B[integrity]     B -- "Min Occurs 1, Max Occurs 1" --&gt; C </pre>
Used by	Element Group specimen.section.content
Model	integrity

Children	integrity
Source	<pre>&lt;xs:group name="specimen.integrity.item"&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.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group specimen.integrity.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element specimen.integrity.item/integrity
Model	response
Children	response
Source	<pre>&lt;xs:group name="specimen.integrity.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="specimen.integrity.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group diagnosis.section.content
Model	tumor
Children	tumor
Source	<pre>&lt;xs:group name="tumor.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="tumor"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="tumor.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.section.content

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element tumor.section/tumor
Model	focus
Children	focus
Source	<pre>&lt;xs:group name="tumor.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group maxOccurs="unbounded" ref="tumor.focus.section"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.focus.section

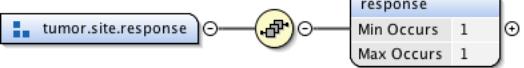
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group tumor.section.content
Model	focus
Children	focus
Source	<pre>&lt;xs:group name="tumor.focus.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="focus"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="tumor.site.item"/&gt;           &lt;xs:group ref="tumor.size.item"/&gt;           &lt;xs:group ref="tumor.histologicType.item"/&gt;           &lt;xs:group ref="tumor.grade.item"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="type" type="tumor.focus.type.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.site.item

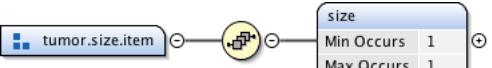
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.focus.section/focus
Model	site
Children	site
Source	<pre>&lt;xs:group name="tumor.site.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="site"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;          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

	</xs:group>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

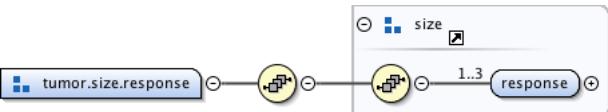
### Element Group tumor.site.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.site.item/site
Model	response
Children	response
Source	<pre>&lt;xs:group name="tumor.site.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="tumor.site.valueset" use="required"/&gt;         &lt;xs:attribute name="on" type="laterality.valueset"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.size.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.focus.section/focus
Model	size
Children	size
Source	<pre>&lt;xs:group name="tumor.size.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="size"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="tumor.size.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.size.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.size.item/size
Model	obbr:response{1,3}
Children	obbr:response
Source	<pre>&lt;xs:group name="tumor.size.response"&gt;   &lt;xs:sequence&gt;</pre>

	<pre> &lt;xs:group ref="size" /&gt; &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.histologicType.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.focus.section/focus
Model	histologicType
Children	histologicType
Source	<pre> &lt;xs:group name="tumor.histologicType.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="histologicType"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content" /&gt;           &lt;xs:group ref="tumor.histologicType.response" /&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.histologicType.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.histologicType.item/histologicType
Model	response
Children	response
Source	<pre> &lt;xs:group name="tumor.histologicType.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.histologicType.valueset" /&gt;           &lt;/xs:simpleType&gt;         &lt;/xs:attribute&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.grade.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.focus.section/focus
Model	grade

Children	grade
Source	<pre>&lt;xs:group name="tumor.grade.item"&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:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group tumor.grade.response

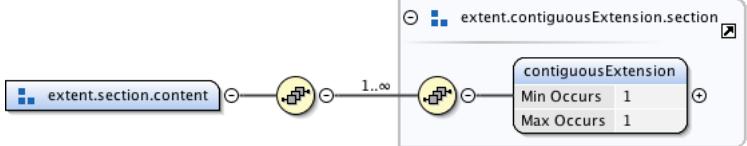
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element tumor.grade.item/grade
Model	response
Children	response
Source	<pre>&lt;xs:group name="tumor.grade.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" type="tumor.grade.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group extent.section

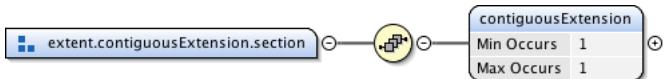
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group diagnosis.section.content
Model	extent
Children	extent
Source	<pre>&lt;xs:group name="extent.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="extent"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="extent.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group extent.section.content

Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element extent.section/extent
Model	contiguousExtension
Children	contiguousExtension
Source	<pre>&lt;xs:group name="extent.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group maxOccurs="unbounded" ref="extent.contiguousExtension.section"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group extent.contiguousExtension.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group extent.section.content
Model	contiguousExtension
Children	contiguousExtension
Source	<pre>&lt;xs:group name="extent.contiguousExtension.section"&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="extent.contiguousExtension.status.section"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group extent.contiguousExtension.status.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element extent.contiguousExtension.section/contiguousExtension
Model	status
Children	status
Source	<pre>&lt;xs:group name="extent.contiguousExtension.status.section"&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 maxOccurs="2" ref="extent.contiguousExtension.status.result.item"/&gt;           &lt;xs:group maxOccurs="2" minOccurs="0" ref="extent.contiguousExtension.status.distance.item"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="of" type="extent.status.of.valueset"/&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>

	</xs:group>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group extent.contiguousExtension.status.result.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[extent.contiguousExtension.status.result.item] --&gt; B(result)     B -- "Min Occurs: 1, Max Occurs: 1" --&gt; C[ ]   </pre>
Used by	Element extent.contiguousExtension.status.section/status
Model	result
Children	result
Source	<pre> &lt;xs:group name="extent.contiguousExtension.status.result.item"&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="extent.contiguousExtension.status.result.response"/&gt;                 &lt;/xs:sequence&gt;                 &lt;xs:attribute name="for" type="extent.contiguousExtension.status.result.for.valueset"/&gt;                 &lt;xs:attribute name="on" type="laterality.valueset"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group extent.contiguousExtension.status.result.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[extent.contiguousExtension.status.result.response] --&gt; B(response)     B -- "Min Occurs: 1, Max Occurs: 1" --&gt; C[ ]   </pre>
Used by	Element extent.contiguousExtension.status.result.item/result
Model	response
Children	response
Source	<pre> &lt;xs:group name="extent.contiguousExtension.status.result.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="qualitative.valueset" use="required"/&gt;                 &lt;xs:attribute name="extent" type="focality.valueset"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group extent.contiguousExtension.status.distance.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[extent.contiguousExtension.status.distance.item] --&gt; B(distance)     B -- "Min Occurs: 1, Max Occurs: 1" --&gt; C[ ]   </pre>
Used by	Element extent.contiguousExtension.status.section/status
Model	distance
Children	distance

Source	<pre>&lt;xs:group name="extent.contiguousExtension.status.distance.item"&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:attribute name="for" type="extent.contiguousExtension.status.distance.for.valueset"/&gt;         &lt;xs:attribute name="from" type="val:base.values"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group obbr:length

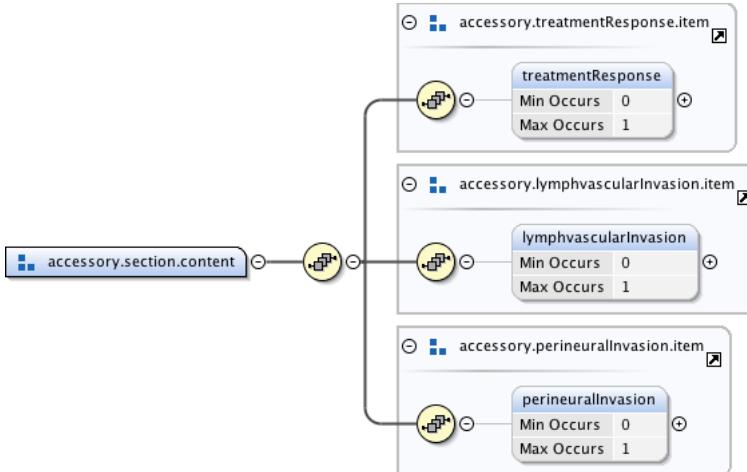
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> classDiagram     class length     class response     length o--&gt; response   </pre>
Used by	Elements extent.contiguousExtension.status.distance.item/distance, margins.margin.distance.item/distance
Model	obbr:response
Children	obbr:response
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="obbr:generic.content"/&gt;         &lt;xs:attributeGroup ref="obbr:quantity"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Element Group accessory.section

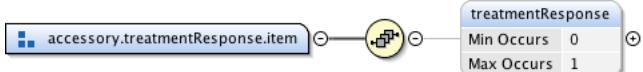
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> classDiagram     class accessory     class diagnosis     class section     class content     class accessorySection     accessorySection o--&gt; diagnosis     accessorySection o--&gt; section     accessorySection o--&gt; content     Min Occurs: 1     Max Occurs: 1   </pre>
Used by	Element Group diagnosis.section.content
Model	accessory
Children	accessory
Source	<pre>&lt;xs:group name="accessory.section"&gt;   &lt;xs:sequence&gt;     &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.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group accessory.section.content

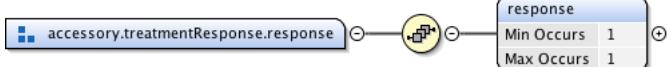
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element accessory.section/accessory
Model	treatmentResponse{0,1} , lymphvascularInvasion{0,1} , perineuralInvasion{0,1}
Children	lymphvascularInvasion, perineuralInvasion, treatmentResponse
Source	<pre>&lt;xs:group name="accessory.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="accessory.treatmentResponse.item"/&gt;     &lt;xs:group ref="accessory.lymphvascularInvasion.item"/&gt;     &lt;xs:group ref="accessory.perineuralInvasion.item"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group accessory.treatmentResponse.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group accessory.section.content
Model	treatmentResponse{0,1}
Children	treatmentResponse
Source	<pre>&lt;xs:group name="accessory.treatmentResponse.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element minOccurs="0" name="treatmentResponse"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="accessory.treatmentResponse.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group accessory.treatmentResponse.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element accessory.treatmentResponse.item/treatmentResponse
Model	response

Children	response
Source	<pre>&lt;xs:group name="accessory.treatmentResponse.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="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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group accessory.lymphvascularInvasion.item

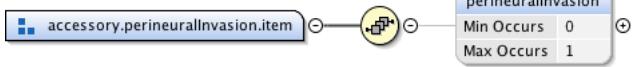
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group accessory.section.content
Model	lymphvascularInvasion{0,1}
Children	lymphvascularInvasion
Source	<pre>&lt;xs:group name="accessory.lymphvascularInvasion.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element minOccurs="0" name="lymphvascularInvasion"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="accessory.lymphvascularInvasion.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group accessory.lymphvascularInvasion.response

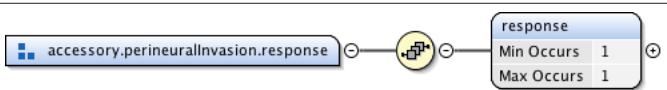
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element accessory.lymphvascularInvasion.item/lymphvascularInvasion
Model	response
Children	response
Source	<pre>&lt;xs:group name="accessory.lymphvascularInvasion.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" type="accessory.lymphvascularInvasion.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group accessory.perineuralInvasion.item

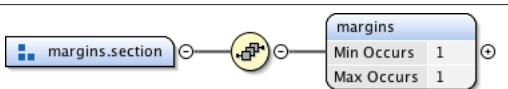
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element Group accessory.section.content
Model	perineuralInvasion{0,1}
Children	perineuralInvasion
Source	<pre>&lt;xs:group name="accessory.perineuralInvasion.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element minOccurs="0" name="perineuralInvasion"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="accessory.perineuralInvasion.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group accessory.perineuralInvasion.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element accessory.perineuralInvasion.item/perineuralInvasion
Model	response
Children	response
Source	<pre>&lt;xs:group name="accessory.perineuralInvasion.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="accessory.perineuralInvasion.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group margins.section

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

Schema location

file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element Group margins.section.content**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[margins.section.content] --&gt; B[margins.margin.section]     B --&gt; C[margin]     C -- "Min Occurs 1, Max Occurs 1" --&gt; D   </pre>
Used by	Element margins.section/margins
Model	margin
Children	margin
Source	<pre> &lt;xs:group name="margins.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group maxOccurs="unbounded" ref="margins.margin.section"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;   </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

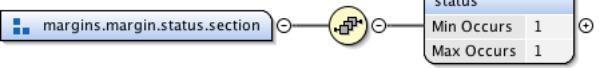
**Element Group margins.margin.section**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[margins.margin.section] --&gt; B[margin]     B -- "Min Occurs 1, Max Occurs 1" --&gt; C   </pre>
Used by	Element Group margins.section.content
Model	margin
Children	margin
Source	<pre> &lt;xs:group name="margins.margin.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="margin"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="margins.margin.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

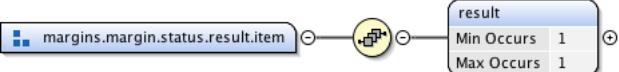
**Element Group margins.margin.section.content**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[margins.margin.section.content] --&gt; B[margins.margin.status.section]     B --&gt; C[status]     C -- "Min Occurs 1, Max Occurs 1" --&gt; D   </pre>
Used by	Element margins.margin.section/margin
Model	status
Children	status
Source	<pre> &lt;xs:group name="margins.margin.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="margins.margin.status.section"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;   </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

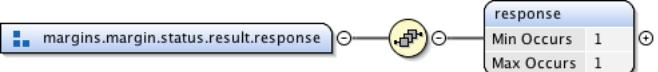
**Element Group margins.margin.status.section**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group margins.margin.section.content
Model	status
Children	status
Source	<pre>&lt;xs:group name="margins.margin.status.section"&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.item"/&gt;           &lt;xs:group minOccurs="0" ref="margins.margin.distance.item"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="of" type="margins.margin.status.of.valueset" use="required"/&gt;         &lt;xs:attribute name="on" type="laterality.valueset"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element Group margins.margin.status.result.item**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element margins.margin.status.section/status
Model	result
Children	result
Source	<pre>&lt;xs:group name="margins.margin.status.result.item"&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="margins.margin.status.result.response"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="for" type="margins.margin.status.result.for.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element Group margins.margin.status.result.response**

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

```

<xs:element name="response">
  <xs:complexType>
    <xs:group ref="generic.content"/>
    <xs:attribute name="value" type="qualitative.valueset" use="required"/>
    <xs:attribute name="extent" type="focality.valueset"/>
    <xs:attribute name="closest" type="qualitative.valueset"/>
    <xs:attributeGroup ref="generic.attrib"/>
  </xs:complexType>
</xs:element>
</xs:sequence>
</xs:group>

```

Schema location file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group margins.margin.distance.item

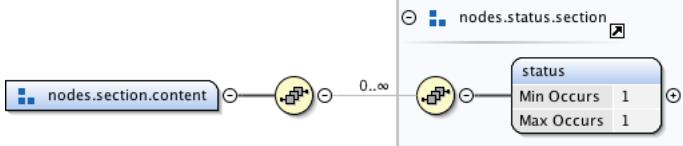
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[margins.margin.distance.item] --- B(( ))     B --- C[distance]     C --- D[Min Occurs: 1]     C --- E[Max Occurs: 1]   </pre>
Used by	Element margins.margin.status.section/status
Model	distance
Children	distance
Source	<pre> &lt;xs:group name="margins.margin.distance.item"&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:attribute name="for" type="margins.margin.distance.for.valueset" use="required"/&gt;         &lt;!--&lt;xs:attribute name="from" type="margins.margin.distance.from.valueset"/--&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group nodes.section

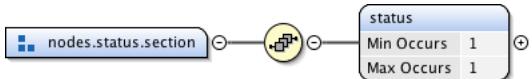
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	<pre> graph LR     A[nodes.section] --- B(( ))     B --- C[nodes]     C --- D[Min Occurs: 1]     C --- E[Max Occurs: 1]   </pre>
Used by	Element Group diagnosis.section.content
Model	nodes
Children	nodes
Source	<pre> &lt;xs:group name="nodes.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="nodes"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="nodes.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group nodes.section.content

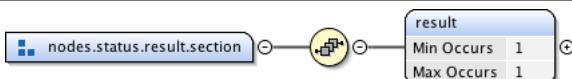
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element nodes.section/nodes
Model	status
Children	status
Source	<pre>&lt;xs:group name="nodes.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="nodes.status.section"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group nodes.status.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group nodes.section.content
Model	status
Children	status
Source	<pre>&lt;xs:group name="nodes.status.section"&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="nodes.status.result.section"/&gt;           &lt;xs:group ref="nodes.status.largestMetastasis.item"/&gt;           &lt;xs:group ref="nodes.status.extracapsularExtension.item"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="of" type="nodes.status.of.valueset" use="required"/&gt;         &lt;xs:attribute name="on" type="laterality.valueset"/&gt;         &lt;xs:attribute name="technique" type="nodes.status.technique.valueset"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group nodes.status.result.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodes.status.section/status
Model	result
Children	result
Source	<pre>&lt;xs:group name="nodes.status.result.section"&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="nodes.status.result.response"/&gt;         &lt;/xs:sequence&gt;         &lt;xs:attribute name="for" type="nodes.status.result.for.valueset" use="required"/&gt;         &lt;xs:attribute name="type" type="nodes.status.result.type.valueset"/&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>

	<pre>         &lt;/xs:element&gt;       &lt;/xs:sequence&gt;     &lt;/xs:group&gt;   </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element Group nodes.status.result.response**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodes.status.result.section/result
Model	response
Children	response
Source	<pre> &lt;xs:group name="nodes.status.result.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:attributeGroup ref="tally"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element Group nodes.status.largestMetastasis.item**

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodes.status.section/status
Model	largestMetastasis
Children	largestMetastasis
Source	<pre> &lt;xs:group name="nodes.status.largestMetastasis.item"&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:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;   </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Element Group nodes.status.extracapsularExtension.item**

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

	<pre> &lt;xs:group ref="generic.content"/&gt; &lt;xs:group ref="nodes.extracapsularExtension.response" /&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group nodes . extracapsularExtension . response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element nodes.status.extracapsularExtension.item/extracapsularExtension
Model	response
Children	response
Source	<pre> &lt;xs:group name="nodes.extracapsularExtension.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="qualitative.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group distant . section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group diagnosis.section.content
Model	distant
Children	distant
Source	<pre> &lt;xs:group name="distant.section"&gt; &lt;xs:sequence&gt; &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.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group distant . section . content

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

Model	status
Children	status
Source	<pre>&lt;xs:group name="distant.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group maxOccurs="unbounded" minOccurs="0" ref="distant.status.section"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group distant.status.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group distant.section.content
Model	status
Children	status
Source	<pre>&lt;xs:group name="distant.status.section"&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="distant.status.section.content"/&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:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group distant.status.section.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element distant.status.section/status
Model	result
Children	result
Source	<pre>&lt;xs:group name="distant.status.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="distant.status.result.item"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group distant.status.result.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group distant.status.section.content
Model	result

Children	result
Source	<pre>&lt;xs:group name="distant.status.result.item"&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:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group distant.status.result.response

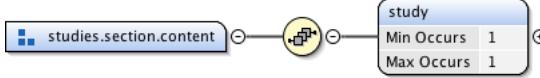
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element distant.status.result.item/result
Model	response
Children	response
Source	<pre>&lt;xs:group name="distant.status.result.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" type="qualitative.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group studies.section

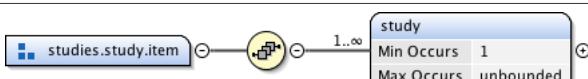
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group diagnosis.section.content
Model	studies
Children	studies
Source	<pre>&lt;xs:group name="studies.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="studies"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="studies.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group studies.section.content

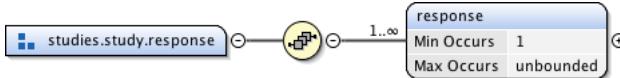
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element studies.section/studies
Model	study
Children	study
Source	<pre>&lt;xs:group name="studies.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="study"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="studies.study.item"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group **studies.study.item**

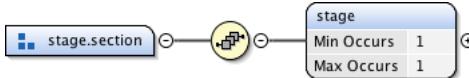
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element studies.section.content/study
Model	study+
Children	study
Source	<pre>&lt;xs:group name="studies.study.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element maxOccurs="unbounded" name="study"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="studies.study.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group **studies.study.response**

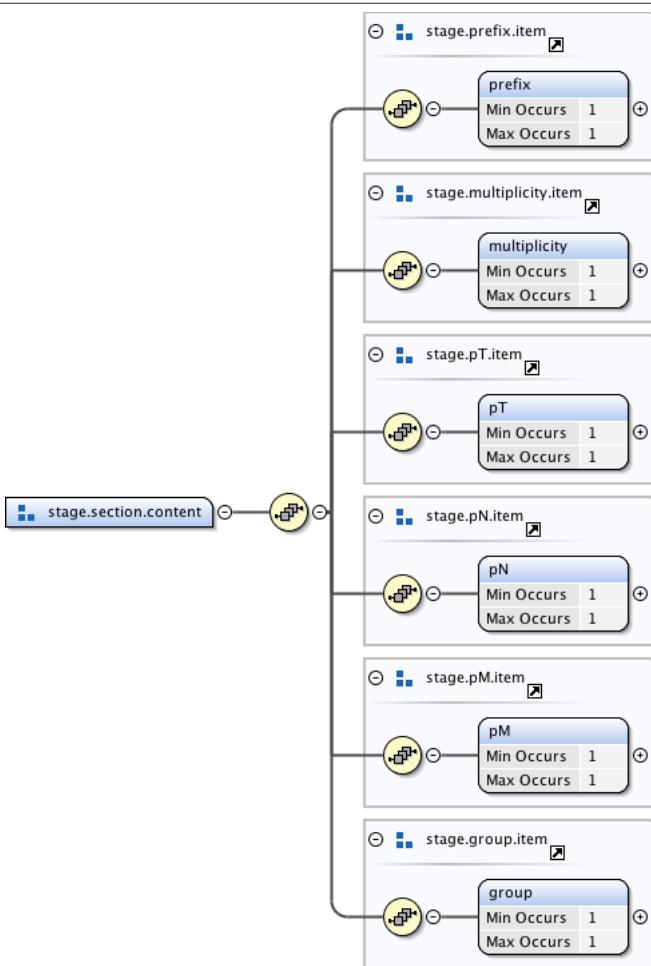
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element studies.study.item/study
Model	response+
Children	response
Source	<pre>&lt;xs:group name="studies.study.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element maxOccurs="unbounded" name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" type="xs:token" 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>

Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd
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## Element Group stage.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group diagnosis.section.content
Model	stage
Children	stage
Source	<pre>&lt;xs:group name="stage.section"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="stage"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="stage.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

## Element Group stage.section.content

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

Model	prefix , multiplicity , pT , pN , pM , group
Children	group, multiplicity, pM, pN, pT, prefix
Source	<pre>&lt;xs:group name="stage.section.content"&gt;   &lt;xs:sequence&gt;     &lt;xs:group ref="stage.prefix.item"/&gt;     &lt;xs:group ref="stage.multiplicity.item"/&gt;     &lt;xs:group ref="stage.pT.item"/&gt;     &lt;xs:group ref="stage.pN.item"/&gt;     &lt;xs:group ref="stage.pM.item"/&gt;     &lt;xs:group ref="stage.group.item"/&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group stage.prefix.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.section.content
Model	prefix
Children	prefix
Source	<pre>&lt;xs:group name="stage.prefix.item"&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:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group stage.prefix.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element stage.prefix.item/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" type="stage.prefix.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group stage.multiplicity.item

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

Used by	Element Group	stage.section.content
Model	multiplicity	
Children	multiplicity	
Source	<pre>&lt;xs:group name="stage.multiplicity.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="multiplicity"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="stage.multiplicity.response"/&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>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

### Element Group stage.multiplicity.response

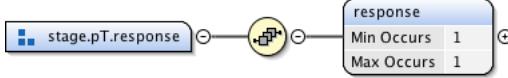
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element
	stage.multiplicity.item/multiplicity
Model	response
Children	response
Source	<pre>&lt;xs:group name="stage.multiplicity.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" type="multiplicity.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group stage.pT.item

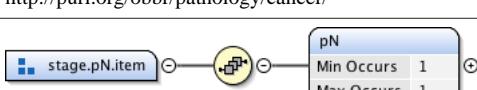
Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group
	stage.section.content
Model	pT
Children	pT
Source	<pre>&lt;xs:group name="stage.pT.item"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="pT"&gt;       &lt;xs:complexType&gt;         &lt;xs:sequence&gt;           &lt;xs:group ref="generic.content"/&gt;           &lt;xs:group ref="stage.pT.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group stage.pT.response

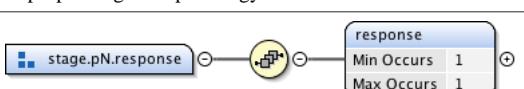
Namespace	http://purl.org/obbr/pathology/cancer/
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Diagram	
Used by	Element stage.pT.item/pT
Model	response
Children	response
Source	<pre>&lt;xs:group name="stage.pT.response"&gt;   &lt;xs:sequence&gt;     &lt;xs:element name="response"&gt;       &lt;xs:complexType&gt;         &lt;xs:group ref="generic.content"/&gt;         &lt;xs:attribute name="value" type="stage.pT.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

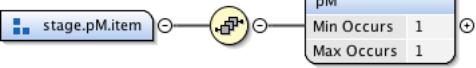
### Element Group stage.pN.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.section.content
Model	pN
Children	pN
Source	<pre>&lt;xs:group name="stage.pN.item"&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.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

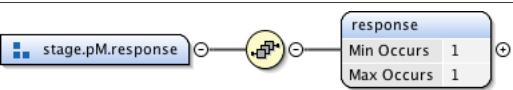
### Element Group stage.pN.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element stage.pN.item/pN
Model	response
Children	response
Source	<pre>&lt;xs:group name="stage.pN.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.pN.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

## Element Group stage.pM.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group stage.section.content
Model	pM
Children	pM
Source	<pre>&lt;xs:group name="stage.pM.item"&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.response"/&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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

## Element Group stage.pM.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element stage.pM.item/pM
Model	response
Children	response
Source	<pre>&lt;xs:group name="stage.pM.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.pM.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

## Element Group stage.group.item

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

	<pre> &lt;/xs:complexType&gt; &lt;/xs:element&gt; &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group stage.group.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element stage.group.item/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:attributeGroup ref="generic.attrib"/&gt;       &lt;/xs:complexType&gt;     &lt;/xs:element&gt;   &lt;/xs:sequence&gt; &lt;/xs:group&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group additional.section

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group diagnosis.section.content
Model	additional
Children	additional
Source	<pre> &lt;xs:group name="additional.section"&gt;   &lt;xs:sequence&gt;     &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.section.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>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group additional.section.content

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element additional.section/additional
Model	finding*
Children	finding

Source	<pre>&lt;xss:group name="additional.section.content"&gt;   &lt;xss:sequence&gt;     &lt;xss:group ref="additional.finding.item" /&gt;   &lt;/xss:sequence&gt; &lt;/xss:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group additional.finding.item

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element Group additional.section.content
Model	finding*
Children	finding
Source	<pre>&lt;xss:group name="additional.finding.item"&gt;   &lt;xss:sequence&gt;     &lt;xss:element maxOccurs="unbounded" minOccurs="0" name="finding"&gt;       &lt;xss:complexType&gt;         &lt;xss:sequence&gt;           &lt;xss:group ref="generic.content"/&gt;           &lt;xss:group ref="additional.finding.response"/&gt;         &lt;/xss:sequence&gt;         &lt;xss:attribute name="of" type="additional.finding.of.valueset"/&gt;         &lt;xss:attributeGroup ref="generic.attrib"/&gt;       &lt;/xss:complexType&gt;     &lt;/xss:element&gt;   &lt;/xss:sequence&gt; &lt;/xss:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

### Element Group additional.finding.response

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Element additional.finding.item/finding
Model	response
Children	response
Source	<pre>&lt;xss:group name="additional.finding.response"&gt;   &lt;xss:sequence&gt;     &lt;xss:element name="response"&gt;       &lt;xss:complexType&gt;         &lt;xss:group ref="generic.content"/&gt;         &lt;xss:attribute name="value" type="additional.finding.valueset" use="required"/&gt;         &lt;xss:attributeGroup ref="generic.attrib"/&gt;       &lt;/xss:complexType&gt;     &lt;/xss:element&gt;   &lt;/xss:sequence&gt; &lt;/xss:group&gt;</pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

## Attribute Groups

### Attribute Group obbr:generic.attrib

Namespace	http://purl.org/obbr/pathology/cancer/
Diagram	
Used by	Elements accessory.lymphvascularInvasion.item/lymphvascularInvasion, accessory.lymphvascularInvasion.response/response, accessory.perineuralInvasion.item/perineuralInvasion, accessory.perineuralInvasion.response/response,

	accessory.section/accessory, accessory.treatmentResponse.item/treatmentResponse, accessory.treatmentResponse.response/response, additional.finding.item/finding, additional.finding.response/response, additional.section/additional, diagnosis, distant.section/distant, distant.status.result.item/result, distant.status.result.response/response, distant.status.section/status, extent.contiguousExtension.section/contiguousExtension, extent.contiguousExtension.status.distance.item/distance, extent.contiguousExtension.status.result.item/result, extent.contiguousExtension.status.result.response/response, extent.contiguousExtension.status.section/status, extent.section/extent, history.clinicalFinding.item/clinicalFinding, history.clinicalFinding.response/response, history.pathologicFinding.item/pathologicFinding, history.pathologicFinding.response/response, history.priorTherapy.item/priorTherapy, history.priorTherapy.response/response, history.radiologicFinding.item/radiologicFinding, history.radiologicFinding.response/response, history.section/history, margins.margin.distance.item/distance, margins.margin.section/margin, margins.margin.status.result.item/result, margins.margin.status.result.response/response, margins.margin.status.section/status, margins.section/margins, nodes.extracapsularExtension.response/response, nodes.section/nodes, nodes.status.extracapsularExtension.item/extracapsularExtension, nodes.status.largestMetastasis.item/largestMetastasis, nodes.status.result.response/response, nodes.status.result.section/result, nodes.status.section/status, specimen.integrity.item/integrity, specimen.integrity.response/response, specimen.procedure.item/procedure, specimen.procedure.response/response, specimen.section/specimen, specimen.site.item/site, specimen.site.response/response, specimen.size.item/size, specimen.weight.item/weight, specimen.weight.response/response, stage.group.item/group, stage.group.response/response, stage.multiplicity.item/multiplicity, stage.multiplicity.response/response, stage.pM.item/pM, stage.pM.response/response, stage.pN.item/pN, stage.pN.response/response, stage.pT.item/pT, stage.pT.response/response, stage.prefix.item/prefix, stage.prefix.response/response, stage.section/stage, studies.section/content/study, studies.section/studies, studies.study.item/study, studies.study.response/response, tumor.focus.section/focus, tumor.grade.item/grade, tumor.grade.response/response, tumor.histologicType.item/histologicType, tumor.histologicType.response/response, tumor.section/tumor, tumor.site.item/site, tumor.site.response/response, tumor.size.item/size				
Attributes	<b>QName</b>	Type	Fixed	Default	Use
	<b>ANY attribute from ANY namespace OTHER than 'http://purl.org/obbr/pathology/cancer/'</b>				
	<b>xml:id</b>	xs:ID			optional
Source	<xs:attributeGroup name="generic.attrib"> <xs:attribute ref="xml:id"/> <xs:anyAttribute namespace="##other" /> </xs:attributeGroup>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd				

### Attribute Group obbr:date

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram	<pre> classDiagram     class date {         @date         @start         @end         @duration     } </pre>				
Used by	Elements distant.status.result.item/result, history.clinicalFinding.item/clinicalFinding, history.pathologicFinding.item/pathologicFinding, history.priorTherapy.item/priorTherapy, history.radiologicFinding.item/radiologicFinding				
Attributes	<b>QName</b>	Type	Fixed	Default	Use
	<b>date</b>	obbr:date.valueset			optional
	<b>duration</b>	xs:duration			optional
	<b>end</b>	obbr:date.valueset			optional
	<b>start</b>	obbr:date.valueset			optional
Source	<xs:attributeGroup name="date">				

	<pre> &lt;xs:attribute name="date" type="obbr:date.valueset"/&gt; &lt;xs:attribute name="start" type="obbr:date.valueset"/&gt; &lt;xs:attribute name="end" type="obbr:date.valueset"/&gt; &lt;xs:attribute name="duration" type="xs:duration"/&gt; &lt;/xs:attributeGroup&gt; </pre>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

### Attribute Group obbr:quantity

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Used by	Elements obbr:length/obbr:response, obbr:size/obbr:response				
Attributes	QName	Type	Fixed	Default	Use
	<b>relation</b>	obbr:relation.valueset			optional
	<b>unit</b>	obbr:lengthUnit.valueset			required
	<b>value</b>	obbr:scalar.valueset			required
Source	<pre> &lt;xs:attributeGroup name="quantity"&gt;   &lt;xs:attribute name="value" type="obbr:scalar.valueset" use="required"/&gt;   &lt;xs:attribute name="unit" type="obbr:lengthUnit.valueset" use="required"/&gt;   &lt;xs:attribute name="relation" type="obbr:relation.valueset"/&gt; &lt;/xs:attributeGroup&gt; </pre>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd				

### Attribute Group obbr:weight

Namespace	http://purl.org/obbr/pathology/cancer/				
Diagram					
Used by	Element specimen.weight.response/response				
Attributes	QName	Type	Fixed	Default	Use
	<b>relation</b>	obbr:relation.valueset			optional
	<b>unit</b>	obbr:weightUnit.valueset			required
	<b>value</b>	obbr:scalar.valueset			required
Source	<pre> &lt;xs:attributeGroup name="weight"&gt;   &lt;xs:attribute name="value" type="obbr:scalar.valueset" use="required"/&gt;   &lt;xs:attribute name="unit" type="obbr:weightUnit.valueset" use="required"/&gt;   &lt;xs:attribute name="relation" type="obbr:relation.valueset"/&gt; &lt;/xs:attributeGroup&gt; </pre>				
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd				

### Attribute Group obbr:tally

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

	<xs:attribute name="relation" type="obbr:relation.valueset"/> </xs:attributeGroup>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

## Namespace: "http://www.w3.org/XML/1998/namespace"

### Schemas

#### Imported schema `xmlid.xsd`

Namespace	http://www.w3.org/XML/1998/namespace
Properties	attribute form default: qualified element form default: unqualified
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/xmlid.xsd

### Attributes

#### Attribute `@xml:id`

Namespace	http://www.w3.org/XML/1998/namespace
Type	xs:ID
Properties	content: simple
Used by	Attribute Group obbr:generic.attrib
Source	<xs:attribute name="id" type="xs:ID"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/xmlid.xsd

## Namespace: ""

### Attributes

#### Attribute `obbr:generic.content / @value`

Namespace	No namespace
Type	obbr:history.clinicalFinding.valueset
Properties	use: required
Used by	Elements accessory.lymphvascularInvasion.response/response, accessory.perineuralInvasion.response/response, accessory.treatmentResponse.response/response, additional.finding.response/response, distant.status.result.response/response, extent.contiguousExtension.status.result.response/response, history.clinicalFinding.response/response, history.pathologicFinding.response/response, history.priorTherapy.response/response, history.radiologicFinding.response/response, margins.margin.status.result.response/response, nodes.extracapsularExtension.response/response, specimen.integrity.response/response, specimen.procedure.response/response, specimen.site.response/response, stage.group.response/response, stage.multiplicity.response/response, stage.pM.response/response, stage.pN.response/response, stage.pT.response/response, stage.prefix.response/response, studies.study.response/response, tumor.grade.response/response, tumor.histologicType.response/response, tumor.site.response/response
Source	<xs:attribute name="value" type="history.clinicalFinding.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

#### Attribute `history.clinicalFinding.response / response / @value`

Namespace	No namespace
Type	obbr:history.clinicalFinding.valueset
Properties	use: required
Source	<xs:attribute name="value" type="history.clinicalFinding.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute obbr:date / @date**

Namespace	No namespace
Type	obbr:date.valueset
Properties	content: simple
Used by	Attribute Group obbr:date
Source	<xs:attribute name="date" type="obbr:date.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:date / @start**

Namespace	No namespace
Type	obbr:date.valueset
Properties	content: simple
Used by	Attribute Group obbr:date
Source	<xs:attribute name="start" type="obbr:date.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:date / @end**

Namespace	No namespace
Type	obbr:date.valueset
Properties	content: simple
Used by	Attribute Group obbr:date
Source	<xs:attribute name="end" type="obbr:date.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:date / @duration**

Namespace	No namespace
Type	xs:duration
Properties	content: simple
Used by	Attribute Group obbr:date
Source	<xs:attribute name="duration" type="xs:duration"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute history.pathologicFinding.response / response / @value**

Namespace	No namespace
Type	obbr:history.pathologicFinding.valueset
Properties	use: required
Source	<xs:attribute name="value" type="history.pathologicFinding.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute history.priorTherapy.response / response / @value**

Namespace	No namespace
Type	obbr:history.priorTherapy.valueset
Properties	use: required
Source	<xs:attribute name="value" type="history.priorTherapy.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute history.radiologicFinding.response / response / @value**

Namespace	No namespace
Type	obbr:history.radiologicFinding.valueset
Properties	use: required
Source	<xs:attribute name="value" type="history.radiologicFinding.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute obbr:generic.content / @on**

Namespace	No namespace
Type	obbr:laterality.valueset
Properties	content: simple
Used by	Elements specimen.procedure.response/response, tumor.site.response/response
Source	<xs:attribute name="on" type="laterality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:laterality.valueset
Properties	content: simple
Source	<xs:attribute name="on" type="laterality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:specimen.procedure.response.valueset
Properties	use: required
Source	<xs:attribute name="value" type="specimen.procedure.response.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute obbr:quantity / @value**

Namespace	No namespace
Type	obbr:scalar.valueset
Properties	use: required
Used by	Attribute Group obbr:quantity
Source	<xs:attribute name="value" type="obbr:scalar.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:quantity / @unit**

Namespace	No namespace
Type	obbr:lengthUnit.valueset
Properties	use: required
Used by	Attribute Group obbr:quantity
Source	<xs:attribute name="unit" type="obbr:lengthUnit.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:quantity / @relation**

Namespace	No namespace
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Type	obbr:relation.valueset
Properties	content: simple
Used by	Attribute Group obbr:quantity
Source	<xs:attribute name="relation" type="obbr:relation.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:weight / @value**

Namespace	No namespace
Type	obbr:scalar.valueset
Properties	use: required
Used by	Attribute Group obbr:weight
Source	<xs:attribute name="value" type="obbr:scalar.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:weight / @unit**

Namespace	No namespace
Type	obbr:weightUnit.valueset
Properties	use: required
Used by	Attribute Group obbr:weight
Source	<xs:attribute name="unit" type="obbr:weightUnit.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

**Attribute obbr:weight / @relation**

Namespace	No namespace
Type	obbr:relation.valueset
Properties	content: simple
Used by	Attribute Group obbr:weight
Source	<xs:attribute name="relation" type="obbr:relation.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd

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

Namespace	No namespace
Type	obbr:specimen.site.valueset
Properties	use: required
Source	<xs:attribute name="value" type="specimen.site.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:specimen.integrity.valueset
Properties	use: required
Source	<xs:attribute name="value" type="specimen.integrity.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:laterality.valueset

Properties	content: simple
Source	<xs:attribute name="on" type="laterality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:tumor.site.valueset
Properties	use: required
Source	<xs:attribute name="value" type="tumor.site.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	union of(obbr:base.values, obbr:tumor.histologicType.valueset)
Properties	use: required
Source	<xs:attribute name="value" use="required"> <xs:simpleType> <xs:union memberTypes="val:base.values tumor.histologicType.valueset"/> </xs:simpleType> </xs:attribute>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:tumor.grade.valueset
Properties	use: required
Source	<xs:attribute name="value" type="tumor.grade.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute tumor.grade.item / grade / @system**

Namespace	No namespace
Type	obbr:tumor.gradeSystem.valueset
Properties	use: required
Used by	Element tumor.grade.item/grade
Source	<xs:attribute name="system" type="tumor.gradeSystem.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute tumor.focus.section / focus / @type**

Namespace	No namespace
Type	obbr:tumor.focus.type.valueset
Properties	use: required
Used by	Element tumor.focus.section/focus
Source	<xs:attribute name="type" type="tumor.focus.type.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute obbr:generic.content / @extent**

Namespace	No namespace
Type	obbr:focality.valueset
Properties	content: simple

Used by	Elements	extent.contiguousExtension.status.result.response/response, margins.margin.status.result.response/response
Source		<xs:attribute name="extent" type="focality.valueset"/>
Schema location		file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:focality.valueset
Properties	content: simple
Source	<xs:attribute name="extent" type="focality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:qualitative.valueset
Properties	use: required
Source	<xs:attribute name="value" type="qualitative.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:extent.contiguousExtension.status.result.for.valueset
Properties	content: simple
Used by	Element extent.contiguousExtension.status.result.item/result
Source	<xs:attribute name="for" type="extent.contiguousExtension.status.result.for.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute extent.contiguousExtension.status.result.item / result / @on**

Namespace	No namespace
Type	obbr:laterality.valueset
Properties	content: simple
Used by	Element extent.contiguousExtension.status.result.item/result
Source	<xs:attribute name="on" type="laterality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:extent.contiguousExtension.status.distance.for.valueset
Properties	content: simple
Used by	Element extent.contiguousExtension.status.distance.item/distance
Source	<xs:attribute name="for" type="extent.contiguousExtension.status.distance.for.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:base.values
Properties	content: simple

Facets	enumeration	specify
	enumeration	inapplicable
	enumeration	unreported
Used by	Element	extent.contiguousExtension.status.distance.item/distance
Source	<xs:attribute name="from" type="val:base.valueset"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute extent.contiguousExtension.status.section / status / @of**

Namespace	No namespace	
Type	obbr:extent.status.of.valueset	
Properties	content: simple	
Used by	Element	extent.contiguousExtension.status.section/status
Source	<xs:attribute name="of" type="extent.status.of.valueset"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute accessory.treatmentResponse.response / response / @value**

Namespace	No namespace	
Type	obbr:accessory.treatmentResponse.valueset	
Properties	use: required	
Source	<xs:attribute name="value" type="accessory.treatmentResponse.valueset" use="required"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute accessory.lymphvascularInvasion.response / response / @value**

Namespace	No namespace	
Type	obbr:accessory.lymphvascularInvasion.valueset	
Properties	use: required	
Source	<xs:attribute name="value" type="accessory.lymphvascularInvasion.valueset" use="required"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute accessory.perineuralInvasion.response / response / @value**

Namespace	No namespace	
Type	obbr:accessory.perineuralInvasion.valueset	
Properties	use: required	
Source	<xs:attribute name="value" type="accessory.perineuralInvasion.valueset" use="required"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute obbr:generic.content / @closest**

Namespace	No namespace	
Type	obbr:qualitative.valueset	
Properties	content: simple	
Used by	Element	margins.margin.status.result.response
Source	<xs:attribute name="closest" type="qualitative.valueset"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

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

Namespace	No namespace
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Type	obbr:qualitative.valueset
Properties	content: simple
Source	<xs:attribute name="closest" type="qualitative.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:focality.valueset
Properties	content: simple
Source	<xs:attribute name="extent" type="focality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:qualitative.valueset
Properties	use: required
Source	<xs:attribute name="value" type="qualitative.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:margins.margin.status.result.for.valueset
Properties	use: required
Used by	Element margins.margin.status.result.item/result
Source	<xs:attribute name="for" type="margins.margin.status.result.for.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute margins.margin.distance.item / distance / @for**

Namespace	No namespace
Type	obbr:margins.margin.distance.for.valueset
Properties	use: required
Used by	Element margins.margin.distance.item/distance
Source	<xs:attribute name="for" type="margins.margin.distance.for.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:margins.margin.status.of.valueset
Properties	use: required
Used by	Element margins.margin.status.section/status
Source	<xs:attribute name="of" type="margins.margin.status.of.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:laterality.valueset
Properties	content: simple

Used by	Element	margins.margin.status.section/status
Source		<xs:attribute name="on" type="laterality.valueset"/>
Schema location		file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute obbr:tally / @value**

Namespace	No namespace	
Type	obbr:tally.valueset	
Properties	use:	required
Used by	Attribute Group	obbr:tally
Source	<xs:attribute name="value" type="obbr:tally.valueset" use="required"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd	

**Attribute obbr:tally / @relation**

Namespace	No namespace	
Type	obbr:relation.valueset	
Properties	content: simple	
Used by	Attribute Group	obbr:tally
Source	<xs:attribute name="relation" type="obbr:relation.valueset"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/generic.xsd	

**Attribute nodes.status.result.section / result / @for**

Namespace	No namespace	
Type	obbr:nodes.status.result.for.valueset	
Properties	use: required	
Used by	Element	nodes.status.result.section/result
Source	<xs:attribute name="for" type="nodes.status.result.for.valueset" use="required"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute nodes.status.result.section / result / @type**

Namespace	No namespace	
Type	obbr:nodes.status.result.type.valueset	
Properties	content: simple	
Used by	Element	nodes.status.result.section/result
Source	<xs:attribute name="type" type="nodes.status.result.type.valueset"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

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

Namespace	No namespace	
Type	obbr:qualitative.valueset	
Properties	use: required	
Source		<xs:attribute name="value" type="qualitative.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute nodes.status.section / status / @of**

Namespace	No namespace	
Type	obbr:nodes.status.of.valueset	

Properties	use:	required
Used by	Element	nodes.status.section/status
Source	<xs:attribute name="of" type="nodes.status.of.valueset" use="required"/>	
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd	

**Attribute nodes.status.section / status / @on**

Namespace	No namespace
Type	obbr:laterality.valueset
Properties	content: simple
Used by	Element nodes.status.section/status
Source	<xs:attribute name="on" type="laterality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute nodes.status.section / status / @technique**

Namespace	No namespace
Type	obbr:nodes.status.technique.valueset
Properties	content: simple
Used by	Element nodes.status.section/status
Source	<xs:attribute name="technique" type="nodes.status.technique.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:qualitative.valueset
Properties	use: required
Source	<xs:attribute name="value" type="qualitative.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:distant.status.result.for.valueset
Properties	content: simple
Used by	Element distant.status.result.item/result
Source	<xs:attribute name="for" type="distant.status.result.for.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:distant.status.of.valueset
Properties	use: required
Used by	Element distant.status.section/status
Source	<xs:attribute name="of" type="distant.status.of.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
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Type	obbr:laterality.valueset
Properties	content: simple
Used by	Element distant.status.section/status
Source	<xs:attribute name="on" type="laterality.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute studies.study.response / response / @value**

Namespace	No namespace
Type	xs:token
Properties	use: required
Source	<xs:attribute name="value" type="xs:token" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:stage.prefix.valueset
Properties	use: required
Source	<xs:attribute name="value" type="stage.prefix.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:multiplicity.valueset
Properties	use: required
Source	<xs:attribute name="value" type="multiplicity.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:stage.pT.valueset
Properties	use: required
Source	<xs:attribute name="value" type="stage.pT.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:stage.pN.valueset
Properties	use: required
Source	<xs:attribute name="value" type="stage.pN.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:stage.pM.valueset
Properties	use: required
Source	<xs:attribute name="value" type="stage.pM.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:stage.group.valueset
Properties	use: required
Source	<xs:attribute name="value" type="stage.group.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:additional.finding.valueset
Properties	use: required
Source	<xs:attribute name="value" type="additional.finding.valueset" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

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

Namespace	No namespace
Type	obbr:additional.finding.of.valueset
Properties	content: simple
Used by	Element additional.finding.item/finding
Source	<xs:attribute name="of" type="additional.finding.of.valueset"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute diagnosis / @schemaLocation**

Namespace	No namespace
Type	xs:anyURI
Properties	use: required
Used by	Element diagnosis
Source	<xs:attribute name="schemaLocation" type="xs:anyURI" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd

**Attribute diagnosis / @version**

Namespace	No namespace
Type	xs:string
Properties	use: required
Used by	Element diagnosis
Source	<xs:attribute name="version" type="xs:string" use="required"/>
Schema location	file:/Users/me/Dropbox/Projects/Synoptic-report/Working/framework/no-specific-site.xsd