

### Sequence Feature Settings

Uniprot

Feature Type	Colour	Configuration	Show
chain			<input checked="" type="checkbox"/>
domain			<input checked="" type="checkbox"/>
metal ion-binding site			<input checked="" type="checkbox"/>
modified residue			<input checked="" type="checkbox"/>
transit peptide			<input checked="" type="checkbox"/>
sequence variant			<input checked="" type="checkbox"/>
strand			<input checked="" type="checkbox"/>
helix			<input checked="" type="checkbox"/>
mutagenesis site			<input checked="" type="checkbox"/>
sequence conflict			<input checked="" type="checkbox"/>

### Display settings for domain features

**Colour**

Simple Colour

**Colour by text**

By text of:

**Graduated Colour**

By range of:

Min value:  Max value:  No value:

**Filters**

Join conditions with  And  Or