

```
Public interface Collection {
```

```
Public void equals ();
```

```
Public void add (); }
```

```
Public interface List {
```

```
Pedido pedido = new Pedido ();
```

```
Abstract List abstractlist = new AbstractList ();
```

```
Public void get (); }
```

```
Public class Pedido {
```

```
Array HensdeLinha []; }
```

```
Public abstract class AbstractList implements List {
```

```
public abstract void equal ();
```

```
public void get () { }
```

```
public void add () { } }
```

```
Public class ArrayList {
```

```
public void get () { }
```

```
public void add () { } }
```

Public class Project {

String name, description;

Source [] sources;

Alignment [] alignment;

Reference Sequence [] reference sequence;

Feature [] features; }

Public class Source {

String name;

Sequence [] sequences; }

Public class Sequence {

String sequence ID, format; }

```
Public class AlignmentMember {
```

```
String referenceMember;
```

```
AlignedSegment [] alignedSegments; }
```

```
Public class AlignedSegment {
```

```
String refStart, refEnd, memberStart, memberEnd; }
```

```
Public class Alignment {
```

```
String name, displayName, description;
```

```
AlignmentMember [] alignmentMembers; }
```

```
Public class ReferenceSequence {
```

```
String name, displayName;
```

```
FeatureLocation [] featureLocations; }
```

```
Public class Feature {
```

```
String name, displayName; }
```

```
Public FeatureLocation {
```

```
FeatureSegment [] featureSegments;
```

```
Variation [] variations; }
```

```
Public class FeatureSegment {
```

```
String refStart, refEnd; }
```

```
Public class Variation {
```

```
String name, displayName, description, scannerModuleName, translationType;
```

```
PatternLocation [] patternLocations; }
```

```
Public class PatternLocation {
```

```
String refStart, refEnd, pattern; }
```



```
Public class Population {  
    public String race, ethnicity, primary-language, language-family; }
```

```
Public class Molecular-sample {  
    public String molecule; }
```

```
Public class Anatomic-location {}
```

```
Public class Panel {  
    public String count-unit, type;  
    public long size;  
    public boolean pooled;  
    Individual[] individuals; }
```

```
Public class Individual {  
    public String father-ID, mother-ID, sex, birth-date;  
    public int death-date; }
```

```
Public class Geographic-location {  
    public double max-longitude, max-latitude, min-longitude, min-latitude; }
```

```
Public Taxon {  
    public String rank, scientific-name; }
```