





```
Enum server_status_choices {
  U [note: "Uploaded reads"]
  V [note: "Uploaded alleles"]
  I [note: "In queue"]
  R [note: "Running reads to alleles"]
  S [note: "Running alleles to MGT"]
  H [note: "Holding"]
  C [note: "Complete"]
  D [note: "To download NCBI reads"]
}
```

```
Enum privacy_status_choices {
  PU [note: "Public"]
  PV [note: "Private"]
}
```

```
Enum assignment_status_choices {
  A [note: "Assigned MGT"]
  F [note: "Failed: contamination"]
  G [note: "Failed: genome quality"]
  S [note: "Failed: serovar incompatibility"]
  E [note: "Error: other"]
}
```

```
Enum continent_choices {
  Africa
  Antarctica
  Oceania
  North_America
  South_America
  Europe
  Asia
}
```

```
Enum orientation_type_choices {
  Forward
  Reverse
}
```

```
Enum locus_type_choices {  
  GN [note: "Gene"]  
  PS [note: "Pseudogene"]  
  IG [note: "Intergenic"]  
  PR [note: "Promoter"]  
  OT [note: "Other"]  
}
```

```
Enum aa_choices {  
  A [note: "Adenine"]  
  C [note: "Cytosine"]  
  G [note: "Guanine"]  
  T [note: "Thymine"]  
  N [note: "Unidentified"]  
}
```