

Modèle relationnel

```
Biosample(sampleID str, sample_type str, env_medium str, isolation_source  
str, host_disease str, geo_loc_name str, collection_date str)
```

```
Assembly(AssemblyAccession str, AssemblyName str, Strain str,  
AssemblyStatus str, Length int, Contigs int, ContigN50 int,  
ScaffoldN50 int, WGS str, BioSampleID str, Coverage int,  
AsmReleaseDate_Genbank str, SubmitterOrganization str, Metagenome  
str)
```

```
Assembly(BioSampleID) reference Biosample(sampleID)
```

```
Genomes(AssemblyAccession str, GenomeID str, Locus str, Genes int,  
CDS int, Proteine int, tRNAs int, rRNAs str, ncRNAs int)
```

```
Genomes(AssemblyAccession) reference Assembly(AssemblyAccession)
```

```
Contigs(GenomeID str, ContigsID str)
```

```
Contigs(GenomeID) reference Genomes(GenomeID)
```

Modèle entité association

