

Ecosystem

- species: vector<Species>
- numberSpecies: int
- numberIndependentSpecies: int
- numberPredatorPrey: int

```

+ Ecosystem()
+ getNumberSpecies(): int const
+ getNumberIndependentSpecies(): int const
+ getNumberPredatorPrey(): int const
+ addSpecies(species: species & const): void
+ addIndependentSpecies(independentSpecies: IndependentSpecies & const): void
+ addPredatorPrey(predatorPrey: PredatorPrey & const): void
    
```

IndependentSpecies

- carryingCapacity: double
- time: double
- growthRate: double

```

+ IndependentSpecies()
+ IndependentSpecies(name: string, time: double, growthRate: double, size: double, carryingCapacity: double, pAlleleFrequency: double, qAlleleFrequency: double)
    
```

```

+ IndependentSpecies(name: string, time: double, growthRate: double, size: double, carryingCapacity: double)
    
```

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+ getCarryingCapacity(): double const
    
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```

+ getTime(): double const
    
```

```

+ getGrowthRate(): double const
    
```

```

+ getPopulationSize(): double
    
```

PredatorPrey

Species

- size: double
- name: string
- pAlleleFrequency: double
- qAlleleFrequency: double

```

+ Species()
+ Species(name: string, size: double, pAlleleFrequency: double, qAlleleFrequency: double)
    
```

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+ getName(): string const
+ getpAlleleFrequency(): double const
+ getqAlleleFrequency(): double const
    
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```

+ findPercentDominant(): double const
    
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+ findNumberDominant(): double const
    
```

```

+ findPercentRecessive(): double const
    
```

```

+ findNumberRecessive(): double const
    
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```

+ findHomozygousDominant(): double const
    
```

```

+ findHomozygousRecessive(): double const
    
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```

+ findHeterozygous(): double const
    
```

```

+ findHomozygousDominantRecessive(): double const
    
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```

+ findHeterozygousRecessive(): double const
    
```

```

+ getsize(): double const
    
```