OLD MODEL

ShakespeareMonkey

DNA: Object

ShakespeareMonkey

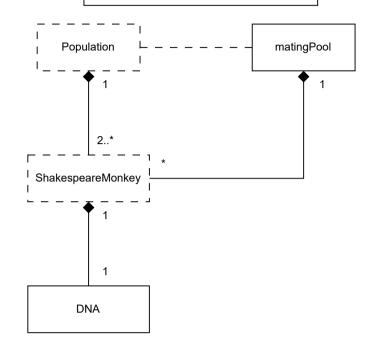
DNA

+ genes: Array
+ fitness: int

+ constructor(int): void
+ getPhrase: String
+ calculateFitness(String): void
+ crossover(DNA): DNA
+ mutate(float): void

related, because the same object comprises them.

But the size of the mating pool is bigger than the population, due to the selection and reproduction steps implementation.



NEW MODEL

ShakespeareMonkey

DNA

+ genes: Array
+ fitness: int

+ constructor(int): void
+ getPhrase: String
+ calculateFitness(String): void
+ crossover(DNA): DNA
+ mutate(float): void

population

+ maxScore: int OR float

+ target: String

+ population: Array

+ generation: int

+ bestFitness: float

+ bestPhrase: String

+ matingPool: Array

+ mutationRate: float

+ finished: bool

+ calculateFitness(): void

+ generateMatingPool(): void

+ getPhrase: String

+ generateNewGenerationAndReplaceOld(): void

+ getBest(): String

+ findMostFitObject(): void

+ trySetFinished(): void

+ isFinished(): bool

+ getGeneratio(): int

+ getAverageFitness(): float

+ getFirstFewPhrases(): String

matingPool is integrated in population.

