Genetic Algorithm: class documentation – Chromosome class

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CLASS NAME		Access Description					
	Data Members		Description				
unsigned char* genes		private	Each array element holds 8 alleles (8 bit data type)				
unsigned int arraySize		private	Holds the size of the array needed to store the whole gene-string (the number of 8-bit elements in the <i>genes</i> array)				
unsigned int fitness		private	Stores the current <i>fitness score</i> assigned to the chromosome				
unsigned int length		private	Length of the gene string (in bits) –all chromosomes in a given				
		<u> </u>	population should have the same number of genes				
Return Value	Function Name		Parameters	Description	Notes		
	default constructor			'			
	custom constructor		unsigned int newlength	The total # of alleles (bits)			
	copy cons		const chromosome& copyChromosome				
	destru	ctor	357511101110001110				
void	AssignFitness		unsigned int newfitness	The new Fitness Score to be assign to the chromosome	The Fitness Score is calculated by a Fitness Function (defined in the main program).		
BOOL (int) 0 = ERROR 1 = SUCCESSFUL COPY	Сору		chromosome* received	pointer to another chromosome object	this chromosome's and the passed chromosome's gene string must be of equal length for the copy to work.		
void	CreateGenes		unsigned int newlength	The length of the gene string (in <i>bits</i> )	Sets up and DMAs the gene array. Assigns a random value to each <i>bit</i> (0 or 1). If default constructor is called then this needs to be called explicitly.		
void	Displa	yFitness			DOS-only		
void	DisplayGenes				DOS-only		
int the decimal value of the extracted portion of the bit	ExtractValue		unsigned int position	bit position of the high order bit that forms the extracted value.	Calculates and returns the decimal value from a defined portion of the gene(bit) string. The returned value is always a positive integer. The maximum value that can be returned is: 32767.		
string (always +)1 signifies that no value could be returned-ERROR.			unsigned int valsize	The # of bits in the extracted portion of the bit string. Max valsize is 15 bits.			
unsigned int Returns the value of fitness (private data member).	GetFit	ness			the Fitness Score of the chromosome.		
unsigned int Returns the value of length (private data member).	GetLe				The bit length of the gene string.		
BOOL (int) 0 = ERROR 1 = SUCCESSFUL INSERTION	Insert	Value	unsigned int position  unsigned int bitlength  unsigned int value	the position of the the high order bit of the inserted value.  The # of bits allocated to representing the inserted value (increased by the function if more bits are required to hold the inserted value).  The value (passed in decimal) to be represented in the	This is the reverse of ExtractValue. The user assigns the position in the bit string where the decimal value is to be inserted and the number of bits allocated to storing this value. The function then attempts to insert the binary equivalent of the decimal value into the bit string (overwriting any bit values occupying the allocated space).		

Return Value	Function Name	Parameters	Description	Notes
int 0 = CAN'T MATE 1 = CROSSOVER 2 = NO	Mate	chromosome* partner	A pointer to the chromosome, which is to be mated with <i>this</i> .	Attempts to mate <i>this</i> chromosome with a selected <i>partner</i> chromosome, which become the 2 <i>children</i>
CROSSOVER		int crossProb	The probability of genetic crossover (the default value of 800 equates to 80%)	chromosomes to be used in the next generation. Crossover probability is passed to the function. Chromosomes must have the same number of genes in order to mate.
void	Mutate	int mutaProb	The probability of mutation of each <i>bit</i> in the gene string. (the default value of 5 equates to 0.5%)	The function moves through each <i>bit</i> of the gene string and decides whether it should be flipped (or mutated) according to the passed probability of mutation.
void	Randomise			Re-randomises the entire gene string of a chromosome (the gene string is initially randomised when it is created).

Genetic Algorithm: class documentation – Population class

CLASS NAME				opaiation diado		
CLASS NAME		Cpopulation				
Data Members		Access	Description			
const unsigned int si	ize	private private	Total number of chromosomes in the population			
	const unsigned int		Length to be assigned to all chromosomes in a population			
chromolen			(default value = 32)			
unsigned int generat		private	Generation counter (incremented by NextGeneration)			
unsigned int mutaPro	ob	private	Mutational Probability to be used in population's mutation phase			
			(default value = 5 (0.5% probability of any gene mutating))			
unsigned int crossPr	ob	private	Cross-over Probability of chromosomes in the population			
			(default value = 800 (80% probability of gene cross-over) )			
chromosome* currer		private	Pointer to first chromosome in current population			
chromosome* mating	gPool	private	Pointer to first chromosome in the selected chromosomes pool			
unsigned int fitness		private	Overall fitness of population (relates to the chromosomes in the			
			currentPool only)			
float meanFitness		private	The average fitness of a chromosome in the population			
unsigned int elitism		private	# of Elite chromosomes to be selected from the <i>current pool</i>			
=			(0 = NO ELITISM)			
unsigned int swapRa	ate	private	(Multi-Point Crossover ONLY) probability of gene swap at the curre			
				locus. (default value = 500 (50% probability of gene swap over) )		
unsigned int eliteCor		private		# of copies to be made of each Elite Chromosome		
Return Value		tion Name	Parameters	Description	Notes	
	defaul	t constructor				
	custor	n constructor	unsigned int	The total # of	All chromosomes in a	
			newsize	chromosomes in	population are created with	
				the population pool	the same number of genes (to	
				(must be an even	allow successful crossover).	
				number)	The population size must be	
			unsigned int newchromolen	The length of a	even so that all chromosomes	
				chromosome's	in the 'mating pool' can be	
				gene-string (in bits)	paired up.	
			unsigned int	The Cross-over		
			newcrossProb	probability		
			unsigned int	The Mutational		
			newmutaProb	Probability		
			unsigned int	# of Elite		
			newelitism	Chromosomes		
			unsigned int	# of copies of Elite		
			neweliteCopies	Chromosomes		
			usigned int	Multi-Point gene		
			newswapRate	crossover rate		
	copy constructor					
	destru				T. 6''	
void	Assig	nFitness	unsigned int	index of the	The Fitness Score is	
			element	chromosome	calculated by a Fitness	
			unsigned int	The new fitness	Function (defined in the main	
			newfitness	value of the	program).	
a sailal	<u> </u>			chromosome	Display data as the	
void	Displa	ayCurrent			Display data on the current	
tol	Diami				population (DOS-only).	
void	ווא	ayMating			Display data on the selection	
					pool (DOS-only).	
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Return Value	Function Name	Parameters	Description	Notes
int the decimal value of the extracted portion of the bit string (always +)1 signifies that no value could be returned-ERROR.	ExtractValue	unsigned int element	The chromosome index in the currentPool array	Calculates and returns the decimal value from a defined portion of the gene(bit) string, of a particular chromosome in the <i>currentPool</i> . The returned value is always a positive integer. The maximum value that can be returned is: 32767.
		unsigned int position  unsigned int	bit position (within the chromosome) of the high order bit that forms the extracted value.  The # of bits in the	
		valsize	extracted portion of the bit string. Max valsize is 15 bits.	
unsigned int Returns the value of chromolen (private data member).	GetChromoLen			Returns the length of chromosomes in the population.
unsigned int Returns the value of crossProb (private data member).	GetcrossProb			Returns the cross-over probability for the population
unsigned int Returns the value of generation (private data member).	GetGeneration			Returns the current generation count
unsigned int Returns the value of mutaProb (private data member).	GetmutaProb			Returns the mutational probability value for chromosomes in the population.
unsigned int Returns the value of size (private data member).	GetSize			Returns the number of chromosomes in the population (this is always an even number and is constant throughout).
BOOL (int) 0 = ERROR 1 = SUCCESSFUL	InsertValue	unsigned int element	The chromosome index in the in the currentPool array	This is the reverse of ExtractValue. Inserts a value into the gene-string of a chromosome.
INSERTION		unsigned int position	the position of the the <i>high order bit</i> of the inserted value (within the chromosome).	
		unsigned int bitlength	The # of bits allocated to representing the inserted value (increased by the function if more bits are required to hold the inserted value).	
		unsigned int value	The value (passed in decimal) to be represented in the allocated portion of the bit string.	

Return Value	Function Name	Parameters	Description	Notes
void	Mate	unsigned int method	0 = Multi-Point Crossover 1 = Single-Point 2 = Two-Point (default = 1)	Mate all chromosomes in selection pool to produce the next generation's (children) chromosomes.
void	Mutate		(doladit 1)	Run the mutation process on each chromosome
void	NextGeneration			Creates the next generation of chromosome objects: copies the matingPool (after mating/mutation has taken place) back into the currentPool, and increment the generation counter.
void	Select	unsigned int method	0 = Roulette Wheel Selection 1 = Tournament Selection	In both selection methods, the fittest chromosomes have the highest probability of being selected for the <i>matingPool</i> . The population's total <i>fitness</i>
		unsigned int sample	# of chromosomes randomly 'sampled' (used in Tournament Selection).	and the fitness of each chromosome (in the current population) needs to have been determined by an external Fitness Function before this is called.