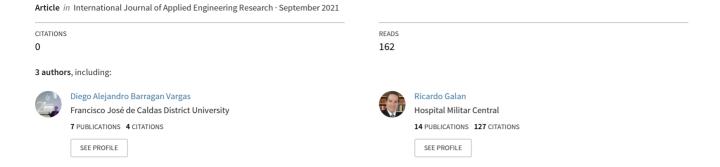
# Design of an Active Band-Pass Filter for the Analysis of Electromyographic Signals Derived from the Median Nerve, Using Genetic Algorithms



## Design of an Active Band-Pass Filter for the Analysis of Electromyographic Signals Derived from the Median Nerve, Using Genetic Algorithms

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#### **Abstract**

In this paper we propose the design of a second order band pass active filter of Sallen Key topology by means of a software developed in MATLAB of a genetic algorithm whit multiobjective optimization, with the purpose of minimizing the sensitivities of the passive components, such as the different resistors and capacitors, in order to reduce the noise and have a measurement signal more faithful to the original, annex to the above, it is intented to minimize the error in the components necessary to filter the signal, with respect to conventional methods that are generally used. As a future work, it is desired to develop a superficial electromyographic device with low sensitivity in its components for the measurement of electrical parameters that are associated with the behavior of the healthy median nerve and with carpal tunnel syndrome, to generate a database and with it different algorithms of analysis of prevention and prediction.

**Keywords:** Band pass filter, operational amplifier, genetic algorithm, sensitivity.

#### 1. Introduction

An electronic filter is a circuit that selects or attenuates a frequency or a range of frequencies, that is, those that pass signals present in certain bands and block signals from other frequency bands, these filters can be divided into two large classes that are known as active filters and passive filters, depending or not on controlled sources and active components, as in the case of operational amplifiers [1]. For the interest of this article, active filters will be analyzed, due to their continuous use in biomedical electronics, since they are necessary for the suppression of noise and amplification of the signal to be studied, it is important to take into account that the implementation of These filters present many options, highlighting designs with topologies of resistors, capacitors and operational amplifiers (active RC filters) [2] [3].

The selection of the discrete components of the RC filters is very important, since the correct operation of the filter in the frequency bands to be attenuated or allowed to pass will depend on this, generally for ease of design equal values are proposed for some components [4-6], this simplifies the process,

however it compromises and limits the freedom of design, which is why the use of genetic algorithms was considered, taking into account that they are heuristic search and optimization methods. In this document we will proceed to design an active band pass filter that takes into account the sensitivities of the components, for this the article [7] that only analyzes an active bi-quadratic low pass filter was taken as a guide. make a comparison with the classic designs that are exposed in the literature to observe the final result of the signal. Filter design is important in biomedical fields such as electromyography, which is a technique used in the functional study of the neuromuscular system, its main objective is the reception of electrical signals produced by a living being [8-10], for Therefore, it is of vital importance to deepen the concept of the devices used and in the same way a review of the different technologies implemented [11], in order to observe in this way the impact of the filters to be designed and with this, take into account factors relevant, such as noise, which greatly affect the desired capture signal [12].

The article was organized as follows: in section 2 the second order high pass filter was shown, in section 3 the second order low pass filter was explained, in section 4 the design of the filters was described with the genetic algorithms, in section 5 the filters were designed in the traditional way, in section 6 the results of the algorithm's behavior and a simulation of the components in MATLAB were shown, making the respective comparison with the traditional design simulation, where detailed the respective information obtained, in section 7 an analysis of the results obtained was carried out and in section 8 the respective conclusions and future work to be carried out were given.

#### 2. Materials and Methods

#### 2.1. High Pass Filter

A high pass filter allows all frequencies above its cutoff frequency to pass through without attenuation [1] [16], in this case a filter with Sallen Key topology will be analyzed as shown in the following figure:

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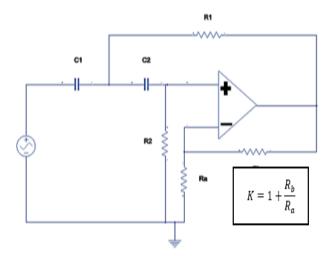


Figure 1. Sallen Key high pass filter

Where:

$$R_a = R \ y \ R_b = (k-1)R$$
 (1)

The transfer function of the high pass filter for this case is:

$$\frac{V_{sal}}{V_{ent}} = \frac{ks^2}{s^2 + \left(\frac{1}{C_1 R_2} + \frac{1}{C_2 R_2} + \frac{(1 - K)}{C_1 R_1}\right)s + \frac{1}{C_1 C_2 R_1 R_2}}$$
(2)

Having:

$$H(s) = \frac{H_0 s^2}{s^2 + \frac{w_n}{O} s + w^2_n} \tag{3}$$

The cutoff frequency is:

$$w_0 = \sqrt{\frac{1}{R_1 R_2 C_1 C_2}} \tag{4}$$

The quality factor is given by:

$$Q = \frac{\sqrt{c_1 c_2 R_1 R_2}}{c_1 R_1 + c_2 R_1 + c_2 R_2 (1 - k)} \tag{5}$$

And the profit is given by:

$$G = \frac{R_2}{R_1} \tag{6}$$

The sensitivity for the filters is:

$$S_{X_i}^F = \frac{dF/F}{dX_i/X_i} \tag{7}$$

Therefore, the sensitivity for the Sallen Key high pass filter is:

$$S_{\kappa}^{W_0} = 0 \tag{8}$$

$$S_{R_1}^{W_0} = S_{R_2}^{W_0} = S_{C_1}^{W_0} = S_{C_2}^{W_0} = -1/2$$
 (9)

$$S_{R_1}^Q = \frac{\sqrt{R_1}}{2} \left( \frac{\sqrt{C_1 C_2 R_2} [C_2 R_2 (1-K) - R_1 (C_1 + C_2)]}{\sqrt{C_1 C_2 R_1 R_2} [R_1 [C_1 + C_2] + C_2 R_2 (1-k)]} \right) (10)$$

$$S_{R_2}^Q = \frac{\sqrt{R_2} \left[ \sqrt{C_1 C_2 R_1} R_1 (C_1 + C_2) - \sqrt{C_1 C_2 R_1} C_2 (1 - k) \right]}{2\sqrt{C_1 C_2 R_1 R_2} \left[ R_1 (C_1 + C_2) + C_2 R_2 (1 - k)^2 \right]} \tag{11}$$

$$S_K^Q = \frac{kC_2R_2}{R_1(C_1 + C_2) + C_2R_2(1 - k)}$$
 (12)

#### 2.2. Low Pass Filter

A Sallen Key low pass filter [17] will be used, as shown in figure 2:

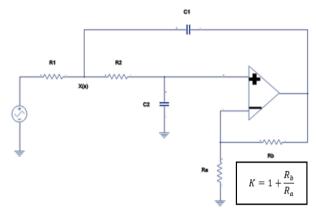


Figure 2. Sallen Key Low Pass Filter

The transfer function for the low pass filter in this case is:

$$\frac{V_{ent}}{V_{sal}} = \frac{\frac{K}{R_1 R_2 C_1 C_2}}{s^2 + \left[\frac{C_2 (R_1 + R_2) + (1 - K) R_1 C_1}{R_1 R_2 C_1 C_2}\right] s + \frac{1}{R_1 R_2 C_1 C_2}}$$
(13)

Having:

$$H(s) = \frac{kw_c^2}{s^2 + \frac{wc}{o}s + w_c^2}$$
 (14)

The cutoff frequency is:

$$w_c = \frac{1}{\sqrt{R_1 R_2 C_1 C_2}} \tag{15}$$

The quality factor is given by:

$$Q = \frac{\sqrt{R_1 R_2 C_1 C_2}}{C_2 (R_1 + R_2) + (1 - k) R_1 C_1} \tag{16}$$

Therefore, the sensitivity for the Sallen Key low pass filter is:

$$S_K^{W_0} = 0 \tag{17}$$

$$S_{R_1}^{W_0} = S_{R_2}^{W_0} = S_{C_1}^{W_0} = S_{C_2}^{W_0} = -1/2$$
 (18)

$$S_{R_1}^Q = \frac{C_2(R_2 - R_1) + C_1(1 - k)(R_1 - 2)}{2[C_2(R_1 + R_2) + R_1C_1(1 - k)]}$$
(19)

$$S_{R_2}^Q = \frac{c_2(R_1 - R_2) + (1 - k)R_1c_1}{c_2(R_1 + R_2) + (1 - k)R_1c_1}$$
 (20)

$$S_K^Q = \frac{R_1 C_1 K}{C_2 (R_1 + R_2) + (1 - k) R_1 C_1} \tag{21}$$

#### 2.3. Filter Design Using Genetic Algorithms

The algorithm must find the values of the passive elements of the filter, since these random values will be set at the beginning to initialize the parameters and therefore the population, to observe and evaluate the sensitivities that depend on the values of the minimum elements and errors in gain, pole frequencies, and quality factor that are less than a specified maximum error. The multiobjective optimization problem that must be solved by the genetic algorithm was proposed in accordance with what was suggested in the article [6], although it differs in the proposed form of the algorithm and in the parameters since a simple genetic algorithm was performed with the primary and secondary parameters, this can be represented by the following expression:

$$minF(y) = S_{R_1}^Q, S_{R_2}^Q, S_{R_3}^Q$$
 (22)

Subject to:

$$Error_G(y) \le Emax$$
 (23)

$$Error_{Wn}(y) \le Emax$$
 (24)

$$Error_{On}(y) \le Emax$$
 (25)

Where Emax is defined as the maximum tolerable design error of the filter characteristics.

Having:

$$Error_{G}(y) = \left| \frac{G(y) - G_{H}}{G_{H}} \right| \tag{26}$$

$$Error_{Wp}(y) = \left| \frac{W_p(y) - W_{pH}}{W_{pH}} \right| \tag{27}$$

$$Error_{Qp}(y) = \left| \frac{Q_p(y) - Q_{pH}}{Q_{pH}} \right| \tag{28}$$

With the conditions of the problem raised, the following steps were taken into account:

• **Population initialization:** It consists of the random generation of an initial population of individuals who are the solutions to the problem [18]. For this case study where the evaluation of a Sallen Key low pass filter and another Sallen Key high pass filter is carried out for the generation of a band pass filter, each individual represents a possible filter configuration characterized by the respective values of resistances and capacitors, where a chromosome made up of five genes is used, as shown below:

Filter High Pass					Filter Low Pass				
R1	R2	K	C1	C2	R1	R2	K	C1	C2

Figure 3 Structure of the generated chromosomes

In the code generated in Matlab, this part was proposed as follows for the two filters:

clear all

close all

clc

%% Population Initialization

inputs = 5; % Number of entries

% Resistors in K\_Ohms

R1=15\*rand(1,1); % Resistor 1

R2=100\*rand(1,1); % Resistor 2

k=20\*rand(1,1); % Resistor 3

C1=0.1\*rand(1,1); % Capacitor 1 in uF

C2=0.2\*rand(1,1); % Capacitor 2 in uF

% Total number of Parameters

totpar = 2\*inputs;

scale=15;

Where the secondary parameters depend on the equations

obtained and for the case of the high pass filter they were:

% Filter secondary parameters

G=R2/R1:

Wp = sqrt(1/(R1\*R2\*C1\*C2));

Qp=(sgrt(C1\*C2\*R1\*R2)/(R1\*(C1+C2)+C2\*R2\*(1-k)));

S1=(sqrt(R1)/2)\*(((sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2)\*((sqrt(C1\*C2\*R2)\*C2\*R2)\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*C2\*R2\*(1-k))-(sqrt(C1\*C2\*R2)\*(1-k))-(sqrt(C1\*C

R2)\*R1\*(C1+C2)))/(sqrt(C1\*C2\*R1\*R2)\*(R1\*(C1+C2)+C 2\*R2\* (1-k))));

S3=((k\*C2\*R2)/(R1\*(C1+C2)+C2\*R2\*(1-k)));

Then the algorithm parameters were generated in code having:

% Genetic Algorithm Parameters

Ngen=500; % Number of Generations

npop=30; % Population size

press=0.01; % Pressure factor selection

ipop=60; % Population mean size

pcross=0.7;% Crossover probability

pmut=0.03;% Mutation Probability

pop=2\*scale\*(rand(npop,totpar)-0.5);

x = [R1, R2, k, C1, C2];

• **Evaluation:** For each generation, the individuals of the present population are evaluated according to a predefined quality criterion that is known as the fitness function, in this article a process similar to that proposed in the article [6] is followed, in which weighted sums methods are used, having:

$$f(y) = (W_1 \left| S_{R_1}^Q \right| + W_2 \left| S_{R_2}^Q \right| + W_3 \left| S_{R_3}^Q \right|) \quad (29)$$

Where the Wi represents the weights assigned to each of the filter sensitivities that must be minimized, in this problem we do not want to give priority to any special weight, so we have:

$$f(y) = \frac{1}{3} (\left| S_{R_1}^Q \right| + \left| S_{R_2}^Q \right| + \left| S_{R_3}^Q \right|) \tag{30}$$

In the Matlab code this part was raised as follows:

% Evolutionary Cycle

for gen =1:Ngen

for i =1:npop% Evaluation of Individuals

% Sensitivity Calculation - Fitness Function

sen=(1/3)\*(abs(S1\*rand(1,1))+abs(S2\*rand(1,1))+abs(S 3\*rand(1,1)));

fobj=sen;

end;

%Subject to:

% Gain error

EG = abs(((G\*sen)-G)/G);

% Pole frequency error

EW = abs(((Wp\*sen)-Wp)/Wp);

% Quality factor error

EQ = abs(((Qp\*sen)-Qp)/Wp);

outpop(gen,1:3)=[ EG EW EQ];

gpop = [ EG EW EQ gen]

- Selection: To generate the next generation of the population, individuals are selected according to their fitness value, for this the roulette method was used, which considers that the probability that an individual is chosen for the crossing is proportional to its fitness value. The roulette selection method was described by Hassoun as "the stochastic version of survival of the fittest", this technique is one of the most used and consists of assigning a segment of the roulette to individuals based on their fitness and the total fitness of the current population and when spinning the roulette as many times as selections are required, the procedure to follow for this is as follows:
- $\circ$  **First:** Calculate the target value f (xi) for each chromosome x (i).
- O **Second:** Calculate the total target value for the population:

$$F = \sum_{i=1}^{I} f(x_i) \ \forall \ i = 1, 2, 3, ..., I$$
 (31)

• Third: Calculate the selection probability pk for each chromosome xi:

$$p_i = \frac{f(x_i)}{F}$$
  $i = 1, 2, 3, ..., I$  (32)

• Fourth: Calculate the cumulative probability pi for each chromosome xi:

$$q_i = \sum_{i=1}^{I} p_i$$
  $i = 1, 2, ..., I$  (33)

```
After the selection, it is done as follows: I repetitions:
                                                                if (pcross>rand(1,1))
                                                                 pind = ceil((totpar-1)*rand(1.1) + 1):
   o First: Generate a random number p in a range [0,1].
                                                                x1 = [indiv1(j,1:pind) indiv2(j,pind+1:totpar)];
   o Second: Choose the i-th chromosome xi such that qi-1
                                                                x2 = [indiv2(j,1:pind) indiv1(j,pind+1:totpar)];
     <p≤ qi.
                                                                indiv1(j,:) = x1;
                                                                indiv2(i,:) = x2;
Carrying out this part in Matlab we have:
                                                                 end
                                                                end
% Qualification of Individuals
Efficiency = (1 - press) * (EG - fobj)/max([EG - EW, eps])
                                                                         Mutation: In this part the uniform random mutation
                                                                operator is used, which considers that each gene has the same
Roulette selection
                                                                probability of being mutated [14] [15]. The code in Matlab
Wheel = cumsum(Efficiency);
                                                                made for this part was:
for j=1:ipop
% Selection of the first couple
                                                                 %Mutation
Shoot = rand(1,1)*max(Wheel);
                                                                 for j=1:ipop
Index = 1:
                                                                 if (pmut>rand(1,1))
while((Wheel(Index)<Shoot)&(Index<length(Wheel)))
                                                                 pind= ceil((totpar-1)*rand(1,1)+1);
Index = Index + 1;
                                                                 vind=2*(rand(1,1)-0.5)*scale;
                                                                 indiv1(j,pind)=vind;
End
indiv1(j,:) = pop(Index,:);
                                                                 end
findiv1(j) = fobj(Index);
                                                                 if (pmut>rand(1,1))
lindiv1(i) = Index;
                                                                 pind= ceil((totpar-1)*rand(1,1)+1);
% Selection of the second partner
                                                                 vind=2*(rand(1,1)-0.5)*scale;
Shoot = rand(1,1)*max(Wheel);
                                                                 indiv2(j,pind)=vind;
Index = 1;
                                                                 end
while((Wheel(Index)<Shoot)&(Index<length(Wheel)))</pre>
                                                                 end
Index = Index + 1;
                                                                 poplast=pop;
end
indiv2(j,:)= pop(Index,:);
                                                                         Stop condition: The cycle is repeated until the stop
                                                                condition is satisfied, which consists of reaching a maximum
findiv2(j) = fobj(Index);
                                                                number of generations. The code generated in Matlab was:
lindiv2(j) = Index;
End
                                                                %New population
                                                                for j=1:npop
        Crossover: The uniform method was used in which
                                                                indexsel= ceil((ipop-1)*rand(1,1)+1);
each gene in the child is obtained by copying the corresponding
gene from one of the parents, according to a randomly
                                                                if rand(1,1) > 0.5
generated binary crossover mask [13] [19]. The code in Matlab
                                                                pop(j,:)=indiv1(indexsel,:);
made for this part was:
                                                                else
                                                                pop(j,:)=indiv2(indexsel,:);
% Crossover
                                                                end
for j=1:ipop
```

end

Wheel = 0:

end

% Obtaining the best individual from the final population

for i =1:npop

w=poplast(i,:);

% Sensitivity calculation Fitness function

sen=(1/3)\*(abs(S1\*rand(1,1))+abs(S2)+abs(S3));

fobi(i,1)=sen;

end

x = x

figure

plot(outpop);

#### 2.4. Filter Design Using the Classic Method

The classical method advises to match some components to facilitate the design. The analysis process generated to obtain the components in the respective filters is shown below, taking into account that the band pass filter will be focused at frequencies from 20Hz to 500Hz.

#### 2.4.1. High-Pass Filter Design

The cutoff frequency for this filter is 20Hz, for ease of design and commercial use 1uF capacitors are used, having C1 = C2, where a quality factor Q = 2 and a resistance R2 =  $20K\Omega$  will be left, having:

$$W_o^2 = \frac{1}{R_1 R_2 C_1 C_2} \tag{34}$$

Solving for R1 we have:

$$R_1 = \frac{1}{R_2 C_1 C_2 W^2_0} = 3,17K\Omega \tag{35}$$

Now we will proceed to clear and find K from equation 4 having:

$$K = 1 - \left[ \frac{\sqrt{C_1 C_2 R_1 R_2} - Q R_1 [C_1 + C_2]}{Q C_2 R_2} \right]$$
 (36)

Where replacing the values you get:

$$K = 1{,}118$$
 (37)

Now we assume that  $Ra = 10K\Omega$  to find Rb, having:

$$R_h = R_a(K - 1) = 2.358K\Omega$$
 (38)

Having with this all the necessary values of the components for the high pass filter.

#### 2.4.2. Low-Pass Filter Design

The cutoff frequency of this filter is 500Hz, for ease of design 0.1uF capacitors are used, an  $R2 = 20K\Omega$  and a quality factor Q = 2 will be used, having:

$$R_1 = \frac{1}{R_2 C_1 C_2 W_0^2} = 506,6\Omega \tag{39}$$

Now we proceed to calculate K having:

$$K = 1 - \frac{1}{R_1 C_1} \left[ \frac{\sqrt{R_1 R_2 C_1 C_2}}{Q} - C_2 (R_1 + R_2) \right]$$
 (40)

Where replacing the values you get:

$$K = 3918.47$$
 (41)

Now we assume that  $Ra = 100\Omega$  to find Rb, having:

$$R_b = R_a(K - 1) = 391,7K\Omega$$
 (42)

Having with this all the necessary values of the components for the low pass filter.

#### 3. Results

#### 3.1. Simulation of Design Filters Using Genetic Algorithms.

It is important to note that the vector x generated in Matlab contains the elements in the following order:

$$x = [R1, R2, k, C1, C2];$$
 (42)

When calculating the respective components with the aforementioned steps, it was obtained:

We proceed to show the simulation of the Sallen Key high pass filter, having:

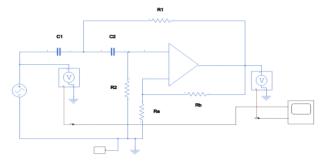


Figure 4. Sallen key high pass filter

Where we proceeded to test with a frequency below 20Hz which is the cutoff frequency, observing that the signal was attenuated for the lower frequencies (in this case it was attenuated 67 times, it was tested with 5Hz) as shown below:

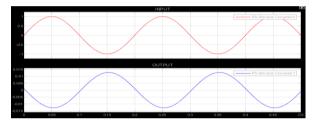


Figure 5. Input signal (Red) and output signal (blue)

Now we proceed to show the values of the vector x obtained for the Sallen Key low-pass filter having:

$$x = [R1, R2, k, C1, C2];$$
 (43)

Having that the values are:

x=0.8749 81.5882 7,3657 0.3208 0.0035

We now proceed to show the simulation for the Sallen Key low pass filter having:

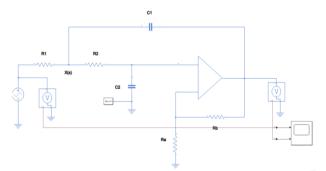


Figure 6. Sallen Key low-pass filter

Where we proceeded to test with a frequency above 500Hz which is the cutoff frequency, observing that the signal was attenuated for higher frequencies as shown below:

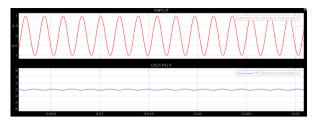


Figure 7. Input signal (Red) and output signal (Blue)

#### 3.2. FILTER SIMULATION CLASSIC DESIGN

We proceed to show the simulation of the Sallen Key high pass filter, having:

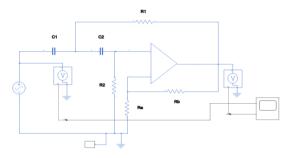


Figure 8. Sallen Key High Pass Filter

Where we proceeded to test with a frequency below 20Hz which is the cutoff frequency, observing that the signal was attenuated for the lower frequencies (in this case it was attenuated 2 times, it was tested with 5Hz) as shown below:

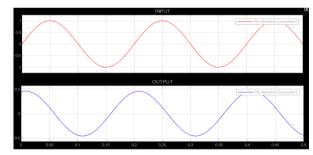


Figure 9. Input signal (Red) and output signal (Blue)

We now proceed to show the simulation for the Sallen Key low pass filter having:

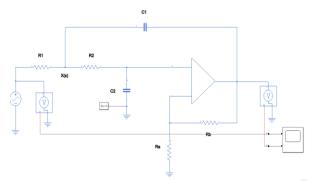


Figure 10. Sallen Key Low Pass Filter

Where we proceeded to test with a frequency above 500Hz which is the cutoff frequency, observing that the signal was attenuated for higher frequencies as shown below:

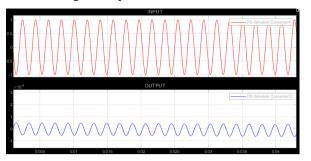


Figure 11. Input signal (Red) and output signal (Blue)

#### 4. Analysis and Discussions Results

After generating the respective calculations to find the components of the designs, the sensitivities of the designs are reviewed having:

#### 4.1. Sensitivity of Filters Designed by Genetic Filter

The vector qpop contains the values of the respective errors and the number of generations iterated to obtain the minimum value of sensitivity.

We proceed to show first the qpop vector of the Sallen Key high-pass filter:

The figure showing the sensitivities of the components is presented below:

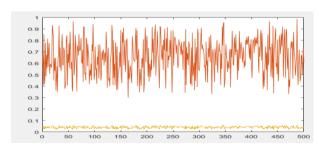


Figure 12. Sensitivity of the resistors in the high pass filter

Now we proceed to show the qpop vector of the Sallen Key low-pass filter, having:

The figure showing the sensitivities of the components is presented below:

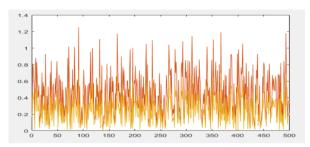


Figure 13. Sensitivities of the resistors in the low-pass filter

### 4.2. SENSITIVITY OF FILTERS DESIGNED IN THE CLASSIC WAY.

We proceed to show the values of the vector qpop, having:

qpop= 0.7395 0.7395 1.2455 500.0000

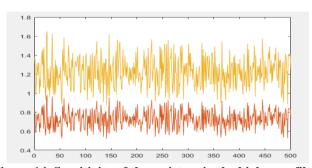


Figure 14. Sensitivity of the resistors in the high pass filter

Now we proceed to show the qpop vector of the Sallen Key low-pass filter, having:

The figure showing the sensitivities of the components is presented below:

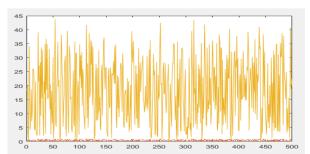


Figure 15. Sensitivity of the resistors in the low pass filter

Carrying out the respective comparison between the sensitivities of the design with genetic algorithms with respect to the classical design, a considerable decrease in the variation of the sensitivity is observed, which is reflected in the errors displayed in the qpop vector, which confirms that the filters have been optimized and These can considerably reduce the noise of the electromyographic signals obtained from the median nerve. A frequency analysis is carried out by means of the Bode diagram.

#### 4.3. High Pass Filter Body Diagram

#### • Classic Design

The transfer function is:

$$H(s) = \frac{1,118s^2}{s^2 + 62,7760s + 15772,87}$$
 (45)

The bode plot of (45) is:

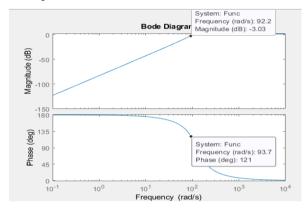


Figure 16. Diagram of the classic design high pass filter winery

#### • Genetic Algorithm

The transfer function is:

$$H(s) = \frac{15,6808s^2}{s^2 - 2221,144s + 11180,135}$$
 (46)

The bode plot of (46) is:

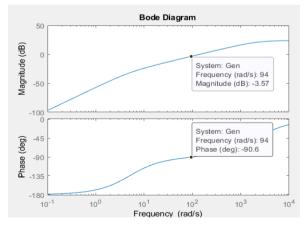


Figure 17. Diagram of the high pass filter design with AG

#### 4.4. Low Pass Filter Body Diagram

#### Classic Design

The transfer function is:

$$H(s) = \frac{378405053,3}{s^2 + 1569,44s + 9869719,7} \tag{47}$$

The bode plot of (47) is:

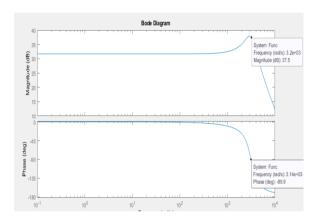


Figure 18. Low pass filter winery diagram classic design

#### • Genetic Algorithm

The transfer function is:

$$H(s) = \frac{91902413,89}{s^2 - 18691,01s + 12477078,06}$$
 (48)

The Bode plot of (48) is:

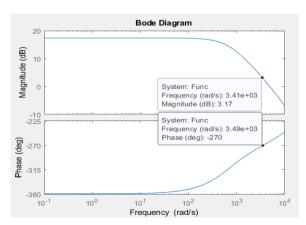


Figure 19. Low pass filter bode diagram AG

#### 5. CONCLUSIONS

- A greater variability was observed in the change in the sensitivity of the filters with the classic design, with respect to the filters of the design with genetic algorithm, this can be evidenced in figures 12 to 15, where the genetic filters have a lower variation, in the case of the genetic high-pass filter, the variation of the sensitivities present in the resistors are between 0 and 16, while in the classic design they are between 0 and 30; In the case of the genetic low-pass filter, the variation in resistive sensitivities is between 1 and 0, while in the classical design it is between 0 and 40.
- The errors obtained with the genetic algorithm are less than those obtained with the classic design, this can be observed in the qpop matrix, because in the genetic design the errors are an average of 10%, while in the classic design the error is found by about 41% on the average of the three errors.
- The Roulette algorithm is sometimes not the most suitable for making the selection since it gives a greater weight to the individual with a greater probability of obtaining suitable characteristics.
- As future work, it is desired to develop a superficial electromyographic device with low sensitivity in its components for the measurement of the electrical parameters that are associated with the behavior of the healthy median nerve and with carpal tunnel syndrome, to generate a database and with it different algorithms of prevention and prediction analysis.

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