

## API Operation

Below are the steps required to retrieve information via the API, with their corresponding java script code snippets. Run these snippets from a browser developer tools console. Note, these snippets can run from a browser, these examples were tested on Chrome version 62.0.3202.94 but should work on any browser.

1. Login to the system and retrieve an access token:

```
var http = new XMLHttpRequest();
var url = "https://api.neurosteer.com/signin";

var params = "email=YOUR_EMAIL&password=YOUR_PASSWORD";

http.open("POST", url, true);

//Send the proper header information along with the request

http.setRequestHeader("Content-type", "application/x-www-form-urlencoded");

http.onreadystatechange = function() {

   if(http.readyState == 4 && http.status == 200) {

      console.log(http.responseText);
   }

}

http.send(params);
```

- 2. To obtain the YOUR ACCESS TOKEN read the response header named access token.
- 3. Open a websocket and listen to updates:



```
// put your business logic here
                          console.log( record.features ); // print list of biomarkers
                 };
                 socket.onopen = function ( event ) {
                          var ta = document.getElementById( 'responseText' );
                           console.log( "Web Socket opened!");
                 };
                 socket.onerror = function () {
                          console.log( 'socket error' );
                           setTimeout( connect, reconnectInterval );
                 };
                 socket.onclose = function ( event ) {
                          console.log( 'socket close' );
                          var ta = document.getElementById( 'responseText' );
                           console.log( "Web Socket closed, reconnect in " + reconnectInterval + " msec" );
                           setTimeout( connect, reconnectInterval );
                 };
        };
        connect();
} else {
        alert( "Your browser does not support Web Socket." );
}
```

4. The Server will respond in a list of biomarkers, updated every second in a JSON format. e.g. {gmean: -0.9990389103528362, c1: -0.9999145017642647, c2: -0.9997699083337485, c3: -0.9999997202373928, h1: -0.9996382954629582, ...}