

```

//a connection between 2 nodes
class connectionGene {
  constructor(from, to, w, inno) {
    this.fromNode = from;
    this.toNode = to;
    this.weight = w;
    this.enabled = true;
    this.innovationNo = inno; //each connection is given a innovation number to compare genomes
  }

  //-----
  //changes the this.weight
  mutateWeight() {
    var rand2 = random(1);
    if (rand2 < 0.1) { //10% of the time completely change the this.weight
      this.weight = random(-1, 1);
    } else { //otherwise slightly change it
      this.weight += (randomGaussian() / 50);
      //keep this.weight between bounds
      if (this.weight > 1) {
        this.weight = 1;
      }
      if (this.weight < -1) {
        this.weight = -1;
      }
    }
  }
}

//-----
//returns a copy of this connectionGene
clone(from, to) {
  var clone = new connectionGene(from, to, this.weight, this.innovationNo);
  clone.enabled = this.enabled;

  return clone;
}
}

```

```

class connectionHistory {
  constructor(from, to, inno, innovationNos) {
    this.fromNode = from;
    this.toNode = to;
    this.innovationNumber = inno;
    this.innovationNumbers = []; //the innovation Numbers from the connections of the genome which first had this mutation
    //this represents the genome and allows us to test if another genome is the same
    //this is before this connection was added
    arrayCopy(innovationNos, this.innovationNumbers); //copy (from, to)
  }

  //-----
  //returns whether the genome matches the original genome and the connection is between the same nodes
  matches(genome, from, to) {
    if (genome.genes.length === this.innovationNumbers.length) { //if the number of connections are different then the genomes aren't the same
      if (from.number === this.fromNode && to.number === this.toNode) {
        //next check if all the innovation numbers match from the genome
        for (var i = 0; i < genome.genes.length; i++) {
          if (!this.innovationNumbers.includes(genome.genes[i].innovationNo)) {
            return false;
          }
        }
        //if reached this far then the innovationNumbers match the genes innovation numbers and the connection is between the same nodes
        //so it does match
        return true;
      }
    }
    return false;
  }
}

```

```

class Genome {
    constructor(inputs, outputs, crossover) {
        this.genes = []; //a list of connections between this.nodes which represent the NN
        this.nodes = [];
        this.inputs = inputs;
        this.outputs = outputs;
        this.layers = 2;
        this.nextNode = 0;
        // this.biasNode;
        this.network = []; //a list of the this.nodes in the order that they need to be considered in the NN
        //create input this.nodes

        if (crossover) {
            return;
        }

        for (var i = 0; i < this.inputs; i++) {
            this.nodes.push(new Node(i));
            this.nextNode++;
            this.nodes[i].layer = 0;
        }

        //create output this.nodes
        for (var i = 0; i < this.outputs; i++) {
            this.nodes.push(new Node(i + this.inputs));
            this.nodes[i + this.inputs].layer = 1;
            this.nextNode++;
        }

        this.nodes.push(new Node(this.nextNode)); //bias node
        this.biasNode = this.nextNode;
        this.nextNode++;
        this.nodes[this.biasNode].layer = 0;
    }

    fullyConnect(innovationHistory) {

        //this will be a new number if no identical genome has mutated in the same

        for (var i = 0; i < this.inputs; i++) {
            for (var j = 0; j < this.outputs; j++) {
                var connectionInnovationNumber = this.getInnovationNumber(innovationHistory, this.nodes[i], this.nodes[this.nodes.length - j - 2]);
                this.genes.push(new connectionGene(this.nodes[i], this.nodes[this.nodes.length - j - 2], random(-1, 1), connectionInnovationNumber));
            }
        }

        var connectionInnovationNumber = this.getInnovationNumber(innovationHistory, this.nodes[this.biasNode], this.nodes[this.nodes.length - 2]);
        this.genes.push(new connectionGene(this.nodes[this.biasNode], this.nodes[this.nodes.length - 2], random(-1, 1), connectionInnovationNumber));

        connectionInnovationNumber = this.getInnovationNumber(innovationHistory, this.nodes[this.biasNode], this.nodes[this.nodes.length - 3]);
        this.genes.push(new connectionGene(this.nodes[this.biasNode], this.nodes[this.nodes.length - 3], random(-1, 1), connectionInnovationNumber));
        //add the connection with a random array

        //changed this so if error here
        this.connectNodes();
    }

    //-----
    //returns the node with a matching number
    //sometimes the this.nodes will not be in order
    getNode(nodeNumber) {
        for (var i = 0; i < this.nodes.length; i++) {
            if (this.nodes[i].number == nodeNumber) {
                return this.nodes[i];
            }
        }
        return null;
    }

    //-----
    //adds the conexions going out of a node to that node so that it can acess the next node during feeding forward
    connectNodes() {

        for (var i = 0; i < this.nodes.length; i++) { //clear the connections
            this.nodes[i].outputConnections = [];
        }

        for (var i = 0; i < this.genes.length; i++) { //for each connectionGene
            this.genes[i].fromNode.outputConnections.push(this.genes[i]); //add it to node
        }
    }

    //-----
    //feeding in input values varo the NN and returning output array
    feedForward(inputValues) {
        //set the outputs of the input this.nodes
        for (var i = 0; i < this.inputs; i++) {
            this.nodes[i].outputValue = inputValues[i];
        }
        this.nodes[this.biasNode].outputValue = 1; //output of bias is 1

        for (var i = 0; i < this.network.length; i++) { //for each node in the network engage it(see node class for what this does)
            this.network[i].engage();
        }

        //the outputs are this.nodes[inputs] to this.nodes [inputs+outputs-1]
        var outs = [];
        for (var i = 0; i < this.outputs; i++) {
            outs[i] = this.nodes[this.inputs + i].outputValue;
        }

        for (var i = 0; i < this.nodes.length; i++) { //reset all the this.nodes for the next feed forward
            this.nodes[i].inputSum = 0;
        }

        return outs;
    }

    //-----
    //sets up the NN as a list of this.nodes in order to be engaged
    generateNetwork() {

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this.connectNodes();
this.network = [];
//for each layer add the node in that layer, since layers cannot connect to themselves there is no need to order the this.nodes within a layer

for (var l = 0; l < this.layers; l++) { //for each layer
  for (var i = 0; i < this.nodes.length; i++) { //for each node
    if (this.nodes[i].layer == l) { //if that node is in that layer
      this.network.push(this.nodes[i]);
    }
  }
}
}

//-----
//mutate the NN by adding a new node
//it does this by picking a random connection and disabling it then 2 new connections are added
//1 between the input node of the disabled connection and the new node
//and the other between the new node and the output of the disabled connection
addNode(innovationHistory) {
  //pick a random connection to create a node between
  if (this.genes.length == 0) {
    this.addConnection(innovationHistory);
    return;
  }
  var randomConnection = floor(random(this.genes.length));

  while (this.genes[randomConnection].fromNode == this.nodes[this.biasNode] && this.genes.length != 1) { //dont disconnect bias
    randomConnection = floor(random(this.genes.length));
  }

  this.genes[randomConnection].enabled = false; //disable it

  var newNodeNo = this.nextNode;
  this.nodes.push(new Node(newNodeNo));
  this.nextNode++;
  //add a new connection to the new node with a weight of 1
  var connectionInnovationNumber = this.getInnovationNumber(innovationHistory, this.genes[randomConnection].fromNode, this.getNode(newNodeNo));
  this.genes.push(new connectionGene(this.genes[randomConnection].fromNode, this.getNode(newNodeNo), 1, connectionInnovationNumber));

  connectionInnovationNumber = this.getInnovationNumber(innovationHistory, this.getNode(newNodeNo), this.genes[randomConnection].toNode);
  //add a new connection from the new node with a weight the same as the disabled connection
  this.genes.push(new connectionGene(this.getNode(newNodeNo), this.genes[randomConnection].toNode, this.genes[randomConnection].weight, connectionInnovationNumber));
  this.getNode(newNodeNo).layer = this.genes[randomConnection].fromNode.layer + 1;

  connectionInnovationNumber = this.getInnovationNumber(innovationHistory, this.nodes[this.biasNode], this.getNode(newNodeNo));
  //connect the bias to the new node with a weight of 0
  this.genes.push(new connectionGene(this.nodes[this.biasNode], this.getNode(newNodeNo), 0, connectionInnovationNumber));

  //if the layer of the new node is equal to the layer of the output node of the old connection then a new layer needs to be created
  //more accurately the layer numbers of all layers equal to or greater than this new node need to be incremented
  if (this.getNode(newNodeNo).layer == this.genes[randomConnection].toNode.layer) {
    for (var i = 0; i < this.nodes.length - 1; i++) { //dont include this newest node
      if (this.nodes[i].layer >= this.getNode(newNodeNo).layer) {
        this.nodes[i].layer++;
      }
    }
    this.layers++;
  }
  this.connectNodes();
}

//-----
//adds a connection between 2 this.nodes which aren't currently connected
addConnection(innovationHistory) {
  //cannot add a connection to a fully connected network
  if (this.fullyConnected()) {
    console.log("connection failed");
    return;
  }

  //get random this.nodes
  var randomNode1 = floor(random(this.nodes.length));
  var randomNode2 = floor(random(this.nodes.length));
  while (this.randomConnectionNodesAreShit(randomNode1, randomNode2)) { //while the random this.nodes are no good
    //get new ones
    randomNode1 = floor(random(this.nodes.length));
    randomNode2 = floor(random(this.nodes.length));
  }
  var temp;
  if (this.nodes[randomNode1].layer > this.nodes[randomNode2].layer) { //if the first random node is after the second then switch
    temp = randomNode2;
    randomNode2 = randomNode1;
    randomNode1 = temp;
  }

  //get the innovation number of the connection
  //this will be a new number if no identical genome has mutated in the same way
  var connectionInnovationNumber = this.getInnovationNumber(innovationHistory, this.nodes[randomNode1], this.nodes[randomNode2]);
  //add the connection with a random array
  this.genes.push(new connectionGene(this.nodes[randomNode1], this.nodes[randomNode2], random(-1, 1), connectionInnovationNumber)); //changed this so if error here
  this.connectNodes();
}

//-----
randomConnectionNodesAreShit(r1, r2) {
  if (this.nodes[r1].layer == this.nodes[r2].layer) return true; // if the this.nodes are in the same layer
  if (this.nodes[r1].isConnectedTo(this.nodes[r2])) return true; //if the this.nodes are already connected

  return false;
}

//-----
//returns the innovation number for the new mutation
//if this mutation has never been seen before then it will be given a new unique innovation number
//if this mutation matches a previous mutation then it will be given the same innovation number as the previous one
getInnovationNumber(innovationHistory, from, to) {
  var isNew = true;
  var connectionInnovationNumber = nextConnectionNo;
  for (var i = 0; i < innovationHistory.length; i++) { //for each previous mutation
    if (innovationHistory[i].matches(this, from, to)) { //if match found
      isNew = false; //its not a new mutation
      connectionInnovationNumber = innovationHistory[i].innovationNumber; //set the innovation number as the innovation number of the match
      break;
    }
  }
}

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if (isNew) { //if the mutation is new then create an arraylist of varegers representing the current state of the genome
    var innoNumbers = [];
    for (var i = 0; i < this.genes.length; i++) { //set the innovation numbers
        innoNumbers.push(this.genes[i].innovationNo);
    }

    //then add this mutation to the innovationHistory
    innovationHistory.push(new connectionHistory(from.number, to.number, connectionInnovationNumber, innoNumbers));
    nextConnectionNo++;
}
return connectionInnovationNumber;
}
}
//-----

//returns whether the network is fully connected or not
fullyConnected() {

    var maxConnections = 0;
    var nodesInLayers = []; //array which stored the amount of this.nodes in each layer
    for (var i = 0; i < this.layers; i++) {
        nodesInLayers[i] = 0;
    }
    //populate array
    for (var i = 0; i < this.nodes.length; i++) {
        nodesInLayers[this.nodes[i].layer] += 1;
    }
    //for each layer the maximum amount of connections is the number in this layer * the number of this.nodes infront of it
    //so lets add the max for each layer together and then we will get the maximum amount of connections in the network
    for (var i = 0; i < this.layers - 1; i++) {
        var nodesInFront = 0;
        for (var j = i + 1; j < this.layers; j++) { //for each layer infront of this layer
            nodesInFront += nodesInLayers[j]; //add up this.nodes
        }
        maxConnections += nodesInLayers[i] * nodesInFront;
    }

    if (maxConnections <= this.genes.length) { //if the number of connections is equal to the max number of connections possible then it is full
        return true;
    }

    return false;
}

//-----
//mutates the genome
mutate(innovationHistory) {
    if (this.genes.length == 0) {
        this.addConnection(innovationHistory);
    }

    var rand1 = random(1);
    if (rand1 < 0.8) { // 80% of the time mutate weights

        for (var i = 0; i < this.genes.length; i++) {
            this.genes[i].mutateWeight();
        }
    }

    //5% of the time add a new connection
    var rand2 = random(1);
    if (rand2 < 0.05) {

        this.addConnection(innovationHistory);
    }

    //1% of the time add a node
    var rand3 = random(1);
    if (rand3 < 0.01) {

        this.addNode(innovationHistory);
    }
}

//-----
//called when this Genome is better than the other parent
crossover(parent2) {
    var child = new Genome(this.inputs, this.outputs, true);
    child.genes = [];
    child.nodes = [];
    child.layers = this.layers;
    child.nextNode = this.nextNode;
    child.biasNodes = this.biasNodes;
    var childGenes = []; // new ArrayList<connectionGene>(); //list of genes to be inherited from the parents
    var isEnabled = []; // new ArrayList<Boolean>();
    //all inherited genes
    for (var i = 0; i < this.genes.length; i++) {
        var setEnabled = true; //is this node in the child going to be enabled

        var parent2gene = this.matchingGene(parent2, this.genes[i].innovationNo);
        if (parent2gene != -1) { //if the genes match
            if (!this.genes[i].enabled || !parent2.genes[parent2gene].enabled) { //if either of the matching genes are disabled

                if (random(1) < 0.75) { //75% of the time disabel the childs gene
                    setEnabled = false;
                }
            }
            var rand = random(1);
            if (rand < 0.5) {
                childGenes.push(this.genes[i]);

                //get gene from this fucker
            } else {
                //get gene from parent2
                childGenes.push(parent2.genes[parent2gene]);
            }
        } else { //disjoint or excess gene
            childGenes.push(this.genes[i]);
            setEnabled = this.genes[i].enabled;
        }
        isEnabled.push(setEnabled);
    }

    //since all excess and disjoint genes are inherited from the more fit parent (this Genome) the childs structure is no different from this parent | with exception of dormant connec
    //so all the this.nodes can be inherited from this parent
    for (var i = 0; i < this.nodes.length; i++) {
        child.nodes.push(this.nodes[i].clone());
    }
}

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}

//clone all the connections so that they connect the childs new this.nodes

for (var i = 0; i < childGenes.length; i++) {
  child.genes.push(childGenes[i].clone(child.getNode(childGenes[i].fromNode.number), child.getNode(childGenes[i].toNode.number)));
  child.genes[i].enabled = isEnabled[i];
}

child.connectNodes();
return child;
}

//-----
//returns whether or not there is a gene matching the input innovation number in the input genome
matchingGene(parent2, innovationNumber) {
  for (var i = 0; i < parent2.genes.length; i++) {
    if (parent2.genes[i].innovationNo == innovationNumber) {
      return i;
    }
  }
  return -1; //no matching gene found
}
//-----
//prints out info about the genome to the console
printGenome() {
  console.log("Prvar genome layers:" + this.layers);
  console.log("bias node: " + this.biasNode);
  console.log("this.nodes");
  for (var i = 0; i < this.nodes.length; i++) {
    console.log(this.nodes[i].number + ",");
  }
  console.log("Genes");
  for (var i = 0; i < this.genes.length; i++) { //for each connectionGene
    console.log("gene " + this.genes[i].innovationNo + "From node " + this.genes[i].fromNode.number + "To node " + this.genes[i].toNode.number +
      "is enabled " + this.genes[i].enabled + "from layer " + this.genes[i].fromNode.layer + "to layer " + this.genes[i].toNode.layer + "weight: " + this.genes[i].weight);
  }
  console.log();
}

//-----
//returns a copy of this genome
clone() {
  var clone = new Genome(this.inputs, this.outputs, true);

  for (var i = 0; i < this.nodes.length; i++) { //copy this.nodes
    clone.nodes.push(this.nodes[i].clone());
  }

  //copy all the connections so that they connect the clone new this.nodes

  for (var i = 0; i < this.genes.length; i++) { //copy genes
    clone.genes.push(this.genes[i].clone(clone.getNode(this.genes[i].fromNode.number), clone.getNode(this.genes[i].toNode.number)));
  }

  clone.layers = this.layers;
  clone.nextNode = this.nextNode;
  clone.biasNode = this.biasNode;
  clone.connectNodes();

  return clone;
}
//-----
//draw the genome on the screen
drawGenome(startX, startY, w, h) {
  //i know its ugly but it works (and is not that important) so I'm not going to mess with it
  var allNodes = []; //new ArrayList<ArrayList<Node>>();
  var nodePoses = []; // new ArrayList<PVector>();
  var nodeNumbers = []; // new ArrayList<Integer>();

  //get the positions on the screen that each node is supposed to be in

  //split the this.nodes varo layers
  for (var i = 0; i < this.layers; i++) {
    var temp = []; // new ArrayList<Node>();
    for (var j = 0; j < this.nodes.length; j++) { //for each node
      if (this.nodes[j].layer == i) { //check if it is in this layer
        temp.push(this.nodes[j]); //add it to this layer
      }
    }
    allNodes.push(temp); //add this layer to all this.nodes
  }

  //for each layer add the position of the node on the screen to the node poses arraylist
  for (var i = 0; i < this.layers; i++) {
    fill(255, 0, 0);
    var x = startX + float((i + 1.0) * w) / float(this.layers + 1.0);
    for (var j = 0; j < allNodes[i].length; j++) { //for the position in the layer
      var y = startY + float((j + 1.0) * h) / float(allNodes[i].length + 1.0);
      nodePoses.push(createVector(x, y));
      nodeNumbers.push(allNodes[i][j].number);
    }
  }

  //draw connections
  stroke(0);
  strokeWeight(2);
  for (var i = 0; i < this.genes.length; i++) {
    if (this.genes[i].enabled) {
      stroke(0);
    } else {
      stroke(100);
    }
    var from;
    var to;
    from = nodePoses[nodeNumbers.indexOf(this.genes[i].fromNode.number)];
    to = nodePoses[nodeNumbers.indexOf(this.genes[i].toNode.number)];
    if (this.genes[i].weight > 0) {
      stroke(255, 0, 0);
    } else {
      stroke(0, 0, 255);
    }
    strokeWeight(map(abs(this.genes[i].weight), 0, 1, 0, 3));
    line(from.x, from.y, to.x, to.y);
  }

  //draw this.nodes last so they appear ontop of the connection lines

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for (var i = 0; i < nodePoses.length; i++) {
  fill(255);
  stroke(0);
  strokeWeight(1);
  ellipse(nodePoses[i].x, nodePoses[i].y, 20, 20);
  textSize(10);
  fill(0);
  textAlign(CENTER, CENTER);
  text(nodeNumbers[i], nodePoses[i].x, nodePoses[i].y);
}

// print out neural network info text
// textAlign(RIGHT);
// fill(255);
// textSize(15);
// noStroke();
// text("car angle", nodePoses[0].x - 20, nodePoses[0].y);
// text("touching ground", nodePoses[1].x - 20, nodePoses[1].y);
// text("angular velocity", nodePoses[2].x - 20, nodePoses[2].y);
// text("Distance to ground", nodePoses[3].x - 20, nodePoses[3].y);
// text("gradient", nodePoses[4].x - 20, nodePoses[4].y);
// text("bias", nodePoses[5].x - 20, nodePoses[5].y);
// textAlign(LEFT);
// text("gas", nodePoses[nodePoses.length - 2].x + 20, nodePoses[nodePoses.length - 2].y);
// text("break", nodePoses[nodePoses.length - 1].x + 20, nodePoses[nodePoses.length - 1].y);

}
}

```

```
<html>
<head>
  <meta charset="UTF-8">
  <script language="javascript" type="text/javascript" src="libraries/p5.js"></script>
  <script language="javascript" src="libraries/p5.dom.js"></script>
  <script language="javascript" src="libraries/p5.sound.js"></script>
  <script language="javascript" src="libraries/Box2d.js"></script>

  <script language="javascript" type="text/javascript" src="sketch.js"></script>
  <script language="javascript" type="text/javascript" src="ConnectionGene.js"></script>
  <script language="javascript" type="text/javascript" src="ConnectionHistory.js"></script>
  <script language="javascript" type="text/javascript" src="Node.js"></script>
  <script language="javascript" type="text/javascript" src="Player.js"></script>
  <script language="javascript" type="text/javascript" src="Population.js"></script>
  <script language="javascript" type="text/javascript" src="Species.js"></script>
  <script language="javascript" type="text/javascript" src="Genome.js"></script>

</head>
<body>

  <div id = "main">
    <h2> Neat template </h2>
    <div id = "canvas">
      </div>
    </div>

    <script>
      //center the canvas
      var tempString = ""+((window.innerWidth*0.9- 1026 - window.innerWidth*0.9*0.08 )/2)+ "px";
      document.getElementById("main").style.marginLeft = tempString;

      window.onresize = function(event) {
        var tempString = ""+((window.innerWidth*0.9- 1026 - window.innerWidth*0.9*0.08 )/2)+ "px";
        document.getElementById("main").style.marginLeft = tempString;
      };

    </script>

  </body>
</html>
```



```

class Node {

    constructor(no) {
        this.number = no;
        this.inputSum = 0; //current sum i.e. before activation
        this.outputValue = 0; //after activation function is applied
        this.outputConnections = []; //new ArrayList<connectionGene>();
        this.layer = 0;
        this.drawPos = createVector();
    }

    //-----
    //the node sends its output to the inputs of the nodes its connected to
    engage() {
        if(this.layer != 0) { //no sigmoid for the inputs and bias
            this.outputValue = this.sigmoid(this.inputSum);
        }

        for(var i = 0; i < this.outputConnections.length; i++) { //for each connection
            if(this.outputConnections[i].enabled) { //dont do shit if not enabled
                this.outputConnections[i].toNode.inputSum += this.outputConnections[i].weight * this.outputValue; //add the weighted output to the sum of the inputs of whatever node this no
            }
        }
    }
    //-----
    //not used
    stepFunction(x) {
        if(x < 0) {
            return 0;
        } else {
            return 1;
        }
    }
    //-----
    //sigmoid activation function
    sigmoid(x) {
        return 1.0 / (1.0 + pow(Math.E, -4.9 * x)); //todo check pow
    }
    //-----
    //returns whether this node connected to the parameter node
    //used when adding a new connection
    isConnectedTo(node) {
        if(node.layer == this.layer) { //nodes in the same this.layer cannot be connected
            return false;
        }

        //you get it
        if(node.layer < this.layer) {
            for(var i = 0; i < node.outputConnections.length; i++) {
                if(node.outputConnections[i].toNode == this) {
                    return true;
                }
            }
        } else {
            for(var i = 0; i < this.outputConnections.length; i++) {
                if(this.outputConnections[i].toNode == node) {
                    return true;
                }
            }
        }
    }

    return false;
}
//-----
//returns a copy of this node
clone() {
    var clone = new Node(this.number);
    clone.layer = this.layer;
    return clone;
}
}

```

[illegible]

```
//this is a template to add NEAT ai to any game
//note //<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<replace
//this means that there is some information specific to the game to input here

var nextConnectionNo = 1000;
var population;
var speed = 60;

var showBest = true; //true if only show the best of the previous generation
var runBest = false; //true if replaying the best ever game
var humanPlaying = false; //true if the user is playing

var humanPlayer;

var showBrain = false;
var showBestEachGen = false;
var upToGen = 0;
var genPlayerTemp; //player

var showNothing = false;

//-----
function setup() {
  window.canvas = createCanvas(1280, 720);
  //<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<replace
  population = new Population(500);
  humanPlayer = new Player();
}
//-----
function draw() {
  drawToScreen();
  if (showBestEachGen) { //show the best of each gen
    showBestPlayersForEachGeneration();
  } else if (humanPlaying) { //if the user is controlling the ship[
    showHumanPlaying();
  } else if (runBest) { // if replaying the best ever game
    showBestEverPlayer();
  } else { //if just evolving normally
    if (population.done()) { //if any players are alive then update them
      population.updateAlive();
    } else { //all dead
      //genetic algorithm
      population.naturalSelection();
    }
  }
}
//-----
function showBestPlayersForEachGeneration() {
  if (genPlayerTemp.dead) { //if current gen player is not dead then update it

    genPlayerTemp.look();
    genPlayerTemp.think();
    genPlayerTemp.update();
    genPlayerTemp.show();
  } else { //if dead move on to the next generation
    upToGen++;
    if (upToGen >= population.genPlayers.length) { //if at the end then return to the start and stop doing it
      upToGen = 0;
      showBestEachGen = false;
    } else { //if not at the end then get the next generation
      genPlayerTemp = population.genPlayers[upToGen].cloneForReplay();
    }
  }
}
//-----
function showHumanPlaying() {
  if (humanPlayer.dead) { //if the player isnt dead then move and show the player based on input
    humanPlayer.look();
    humanPlayer.update();
    humanPlayer.show();
  } else { //once done return to ai
    humanPlaying = false;
  }
}
//-----
function showBestEverPlayer() {
  if (population.bestPlayer.dead) { //if best player is not dead
    population.bestPlayer.look();
    population.bestPlayer.think();
    population.bestPlayer.update();
    population.bestPlayer.show();
  } else { //once dead
    runBest = false; //stop replaying it
    population.bestPlayer = population.bestPlayer.cloneForReplay(); //reset the best player so it can play again
  }
}
//-----
//draws the display screen
function drawToScreen() {
  if (showNothing) {
    //pretty stuff
    //<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<replace
    drawBrain();
    writeInfo();
  }
}
//-----
function drawBrain() { //show the brain of whatever genome is currently showing
  var startX = 0; //<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<<replace
  var startY = 0;
  var w = 0;
  var h = 0;

  if (runBest) {
    population.bestPlayer.brain.drawGenome(startX, startY, w, h);
  } else
  if (humanPlaying) {
    showBrain = false;
  } else if (showBestEachGen) {
    genPlayerTemp.brain.drawGenome(startX, startY, w, h);
  }
}
```

```
} else {  
    population.players[0].brain.drawGenome(startX, startY, w, h);  
}  
  
//-----  
//writes info about the current player  
function writeInfo() {  
    fill(200);  
    textAlign(LEFT);  
    textSize(30);  
    if (showBestEachGen) {  
        text("Score: " + genPlayerTemp.score, 650, 50); //<<<<<<<<<<<<<<<<<<<<<<replace  
        text("Gen: " + (genPlayerTemp.gen + 1), 1150, 50);  
    } else  
    if (humanPlaying) {  
        text("Score: " + humanPlayer.score, 650, 50); //<<<<<<<<<<<<<<<<<<<<<<replace  
    } else  
    if (runBest) {  
        text("Score: " + population.bestPlayer.score, 650, 50); //<<<<<<<<<<<<<<<<<<<<<<replace  
        text("Gen: " + population.gen, 1150, 50);  
    } else {  
        if (showBest) {  
            text("Score: " + population.players[0].score, 650, 50); //<<<<<<<<<<<<<<<<<<<<<<replace  
            text("Gen: " + population.gen, 1150, 50);  
            text("Species: " + population.species.length, 50, canvas.height / 2 + 300);  
            text("Global Best Score: " + population.bestScore, 50, canvas.height / 2 + 200);  
        }  
    }  
}  
  
//-----  
  
function keyPressed() {  
    switch (key) {  
        case ' ':  
            //toggle showBest  
            showBest = !showBest;  
            break;  
        // case '+': //speed up frame rate  
        //     speed += 10;  
        //     frameRate(speed);  
        //     prvarln(speed);  
        //     break;  
        // case '-': //slow down frame rate  
        //     if(speed > 10) {  
        //         speed -= 10;  
        //         frameRate(speed);  
        //         prvarln(speed);  
        //     }  
        //     break;  
        case 'B': //run the best  
            runBest = !runBest;  
            break;  
        case 'G': //show generations  
            showBestEachGen = !showBestEachGen;  
            upToGen = 0;  
            genPlayerTemp = population.genPlayers[upToGen].clone();  
            break;  
        case 'N': //show absolutely nothing in order to speed up computation  
            showNothing = !showNothing;  
            break;  
        case 'P': //play  
            humanPlaying = !humanPlaying;  
            humanPlayer = new Player();  
            break;  
    }  
    //any of the arrow keys  
    switch (keyCode) {  
        case UP_ARROW: //the only time up/ down / left is used is to control the player  
            //<<<<<<<<<<<<<<<<<<<<<<replace  
            break;  
        case DOWN_ARROW:  
            //<<<<<<<<<<<<<<<<<<<<<<replace  
            break;  
        case LEFT_ARROW:  
            //<<<<<<<<<<<<<<<<<<<<<<replace  
            break;  
        case RIGHT_ARROW: //right is used to move through the generations  
  
            if (showBestEachGen) { //if showing the best player each generation then move on to the next generation  
                upToGen++;  
                if (upToGen >= population.genPlayers.length) { //if reached the current generation then exit out of the showing generations mode  
                    showBestEachGen = false;  
                } else {  
                    genPlayerTemp = population.genPlayers[upToGen].cloneForReplay();  
                }  
            } else if (humanPlaying) { //if the user is playing then move player right  
  
                //<<<<<<<<<<<<<<<<<<<<<<replace  
            }  
            break;  
    }  
}
```

```
# NEAT Template JavaScript
```

```

class Species {

  constructor(p) {
    this.players = [];
    this.bestFitness = 0;
    this.champ;
    this.averageFitness = 0;
    this.staleness = 0; //how many generations the species has gone without an improvement
    this.rep;

    //-----
    //coefficients for testing compatibility
    this.excessCoeff = 1;
    this.weightDiffCoeff = 0.5;
    this.compatibilityThreshold = 3;
    if (p) {
      this.players.push(p);
      //since it is the only one in the species it is by default the best
      this.bestFitness = p.fitness;
      this.rep = p.brain.clone();
      this.champ = p.cloneForReplay();
    }
  }

  //-----
  //returns whether the parameter genome is in this species
  sameSpecies(g) {
    var compatibility;
    var excessAndDisjoint = this.getExcessDisjoint(g, this.rep); //get the number of excess and disjoint genes between this player and the current species this.rep
    var averageWeightDiff = this.averageWeightDiff(g, this.rep); //get the average weight difference between matching genes

    var largeGenomeNormaliser = g.genes.length - 20;
    if (largeGenomeNormaliser < 1) {
      largeGenomeNormaliser = 1;
    }

    compatibility = (this.excessCoeff * excessAndDisjoint / largeGenomeNormaliser) + (this.weightDiffCoeff * averageWeightDiff); //compatibility formula
    return (this.compatibilityThreshold > compatibility);
  }

  //-----
  //add a player to the species
  addToSpecies(p) {
    this.players.push(p);
  }

  //-----
  //returns the number of excess and disjoint genes between the 2 input genomes
  //i.e. returns the number of genes which dont match
  getExcessDisjoint(brain1, brain2) {
    var matching = 0.0;
    for (var i = 0; i < brain1.genes.length; i++) {
      for (var j = 0; j < brain2.genes.length; j++) {
        if (brain1.genes[i].innovationNo == brain2.genes[j].innovationNo) {
          matching++;
          break;
        }
      }
    }
    return (brain1.genes.length + brain2.genes.length - 2 * (matching)); //return no of excess and disjoint genes
  }

  //-----
  //returns the average weight difference between matching genes in the input genomes
  averageWeightDiff(brain1, brain2) {
    if (brain1.genes.length == 0 || brain2.genes.length == 0) {
      return 0;
    }

    var matching = 0;
    var totalDiff = 0;
    for (var i = 0; i < brain1.genes.length; i++) {
      for (var j = 0; j < brain2.genes.length; j++) {
        if (brain1.genes[i].innovationNo == brain2.genes[j].innovationNo) {
          matching++;
          totalDiff += abs(brain1.genes[i].weight - brain2.genes[j].weight);
          break;
        }
      }
    }
    if (matching == 0) { //divide by 0 error
      return 100;
    }
    return totalDiff / matching;
  }

  //-----
  //sorts the species by fitness
  sortSpecies() {
    var temp = []; // new ArrayList < Player > ();

    //selection short
    for (var i = 0; i < this.players.length; i++) {
      var max = 0;
      var maxIndex = 0;
      for (var j = 0; j < this.players.length; j++) {
        if (this.players[j].fitness > max) {
          max = this.players[j].fitness;
          maxIndex = j;
        }
      }
      temp.push(this.players[maxIndex]);

      this.players.splice(maxIndex, 1);
      // this.players.remove(maxIndex);
      i--;
    }

    // this.players = (ArrayList) temp.clone();
    arrayCopy(temp, this.players);
    if (this.players.length == 0) {
      this.staleness = 200;
      return;
    }
    //if new best player
    if (this.players[0].fitness > this.bestFitness) {
      this.staleness = 0;
      this.bestFitness = this.players[0].fitness;
      this.rep = this.players[0].brain.clone();
    }
  }
}

```

```

        this.champ = this.players[0].cloneForReplay();
    } else { //if no new best player
        this.staleness++;
    }
}

//-----
//simple stuff
setAverage() {
    var sum = 0;
    for (var i = 0; i < this.players.length; i++) {
        sum += this.players[i].fitness;
    }
    this.averageFitness = sum / this.players.length;
}
//-----

//gets baby from the this.players in this species
giveMeBaby(innovationHistory) {
    var baby;
    if (random(1) < 0.25) { //25% of the time there is no crossover and the child is simply a clone of a random(ish) player
        baby = this.selectPlayer().clone();
    } else { //75% of the time do crossover

        //get 2 random(ish) parents
        var parent1 = this.selectPlayer();
        var parent2 = this.selectPlayer();

        //the crossover function expects the highest fitness parent to be the object and the lowest as the argument
        if (parent1.fitness < parent2.fitness) {
            baby = parent2.crossover(parent1);
        } else {
            baby = parent1.crossover(parent2);
        }
    }
    baby.brain.mutate(innovationHistory); //mutate that baby brain
    return baby;
}

//-----
//selects a player based on it fitness
selectPlayer() {
    var fitnessSum = 0;
    for (var i = 0; i < this.players.length; i++) {
        fitnessSum += this.players[i].fitness;
    }
    var rand = random(fitnessSum);
    var runningSum = 0;

    for (var i = 0; i < this.players.length; i++) {
        runningSum += this.players[i].fitness;
        if (runningSum > rand) {
            return this.players[i];
        }
    }
    //unreachable code to make the parser happy
    return this.players[0];
}
//-----
//kills off bottom half of the species
cull() {
    if (this.players.length > 2) {
        for (var i = this.players.length / 2; i < this.players.length; i++) {
            // this.players.remove(i);
            this.players.splice(i, 1);
            i--;
        }
    }
}
//-----
//in order to protect unique this.players, the fitnesses of each player is divided by the number of this.players in the species that that player belongs to
fitnessSharing() {
    for (var i = 0; i < this.players.length; i++) {
        this.players[i].fitness /= this.players.length;
    }
}
}

```

```

class Population {
  constructor(size) {
    this.players = []; //new ArrayList<Player>();
    this.bestPlayer; //the best ever player
    this.bestScore = 0; //the score of the best ever player
    this.globalBestScore = 0;
    this.gen = 1;
    this.innovationHistory = []; // new ArrayList<connectionHistory>();
    this.genPlayers = []; //new ArrayList<Player>();
    this.species = []; //new ArrayList<Species>();

    this.massExtinctionEvent = false;
    this.newStage = false;

    for (var i = 0; i < size; i++) {
      this.players.push(new Player());
      this.players[this.players.length - 1].brain.mutate(this.innovationHistory);
      this.players[this.players.length - 1].brain.generateNetwork();
    }
  }

  updateAlive() {
    for (var i = 0; i < this.players.length; i++) {
      if (!this.players[i].dead) {
        this.players[i].look(); //get inputs for brain
        this.players[i].think(); //use outputs from neural network
        this.players[i].update(); //move the player according to the outputs from the neural network
        if (!showNothing || (showBest || i == 0)) {
          this.players[i].show();
        }
        if (this.players[i].score > this.globalBestScore) {
          this.globalBestScore = this.players[i].score;
        }
      }
    }
  }

  //-----
  //returns true if all the players are dead sad
  done() {
    for (var i = 0; i < this.players.length; i++) {
      if (!this.players[i].dead) {
        return false;
      }
    }
    return true;
  }

  //-----
  //sets the best player globally and for this this.gen
  setBestPlayer() {
    var tempBest = this.species[0].players[0];
    tempBest.gen = this.gen;

    //if best this this.gen is better than the global best score then set the global best as the best this this.gen

    if (tempBest.score >= this.bestScore) {
      this.genPlayers.push(tempBest.cloneForReplay());
      console.log("old best: " + this.bestScore);
      console.log("new best: " + tempBest.score);
      this.bestScore = tempBest.score;
      this.bestPlayer = tempBest.cloneForReplay();
    }
  }

  //-----
  //this function is called when all the players in the this.players are dead and a new this.generation needs to be made
  naturalSelection() {
    // this.batchNo = 0;
    var previousBest = this.players[0];
    this.speciate(); //separate the this.players varo this.species
    this.calculateFitness(); //calculate the fitness of each player
    this.sortSpecies(); //sort the this.species to be ranked in fitness order, best first
    if (this.massExtinctionEvent) {
      this.massExtinction();
      this.massExtinctionEvent = false;
    }
    this.cullSpecies(); //kill off the bottom half of each this.species
    this.setBestPlayer(); //save the best player of this this.gen
    this.killStaleSpecies(); //remove this.species which haven't improved in the last 15(ish) this.generations
    this.killBadSpecies(); //kill this.species which are so bad that they cant reproduce

    // if (this.gensSinceNewWorld >= 0 || this.bestScore > (grounds[0].distance - 350) / 10) {
    //   this.gensSinceNewWorld = 0;
    //   console.log(this.gensSinceNewWorld);
    //   console.log(this.bestScore);
    //   console.log(grounds[0].distance);
    //   newWorlds();
    // }

    console.log("generation " + this.gen + " Number of mutations " + this.innovationHistory.length + " species: " + this.species.length + " <-----");

    var averageSum = this.getAvgFitnessSum();
    var children = [];
    for (var j = 0; j < this.species.length; j++) { //for each this.species
      children.push(this.species[j].champ.clone()); //add champion without any mutation
      var NoOfChildren = floor(this.species[j].averageFitness / averageSum * this.players.length) - 1; //the number of children this this.species is allowed, note -1 is because the c
      for (var i = 0; i < NoOfChildren; i++) { //get the calculated amount of children from this this.species
        children.push(this.species[j].giveMeBaby(this.innovationHistory));
      }
    }
    if (children.length < this.players.length) {
      children.push(previousBest.clone());
    }
    while (children.length < this.players.length) { //if not enough babies (due to flooring the number of children to get a whole var)
      children.push(this.species[0].giveMeBaby(this.innovationHistory)); //get babies from the best this this.gen
    }

    this.players = [];
    arrayCopy(children, this.players); //set the children as the current this.playersulation
    this.gen += 1;
    for (var i = 0; i < this.players.length; i++) { //generate networks for each of the children
      this.players[i].brain.generateNetwork();
    }
  }

  //-----
  //separate this.players into this.species based on how similar they are to the leaders of each this.species in the previous this.gen

```



```

speciate() {
    for (var s of this.species) { //empty this.species
        s.players = [];
    }
    for (var i = 0; i < this.players.length; i++) { //for each player
        var speciesFound = false;
        for (var s of this.species) { //for each this.species
            if (s.sameSpecies(this.players[i].brain)) { //if the player is similar enough to be considered in the same this.species
                s.addToSpecies(this.players[i]); //add it to the this.species
                speciesFound = true;
                break;
            }
        }
        if (!speciesFound) { //if no this.species was similar enough then add a new this.species with this as its champion
            this.species.push(new Species(this.players[i]));
        }
    }
}

//-----
//calculates the fitness of all of the players
calculateFitness() {
    for (var i = 1; i < this.players.length; i++) {
        this.players[i].calculateFitness();
    }
}

//-----
//sorts the players within a this.species and the this.species by their fitnesses
sortSpecies() {
    //sort the players within a this.species
    for (var s of this.species) {
        s.sortSpecies();
    }

    //sort the this.species by the fitness of its best player
    //using selection sort like a loser
    var temp = []; //new ArrayList<Species>();
    for (var i = 0; i < this.species.length; i++) {
        var max = 0;
        var maxIndex = 0;
        for (var j = 0; j < this.species.length; j++) {
            if (this.species[j].bestFitness > max) {
                max = this.species[j].bestFitness;
                maxIndex = j;
            }
        }
        temp.push(this.species[maxIndex]);
        this.species.splice(maxIndex, 1);
        // this.species.remove(maxIndex);
        i--;
    }
    this.species = [];
    arrayCopy(temp, this.species);
}

//-----
//kills all this.species which haven't improved in 15this.generations
killStaleSpecies() {
    for (var i = 2; i < this.species.length; i++) {
        if (this.species[i].staleness >= 15) {
            // .remove(i);
            // splice(this.species, i)
            this.species.splice(i, 1);
            i--;
        }
    }
}

//-----
//if a this.species sucks so much that it wont even be allocated 1 child for the nextthis.generation then kill it now
killBadSpecies() {
    var averageSum = this.getAvgFitnessSum();

    for (var i = 1; i < this.species.length; i++) {
        if (this.species[i].averageFitness / averageSum * this.players.length < 1) { //if wont be given a single child
            // this.species.remove(i); //sad
            this.species.splice(i, 1);
            i--;
        }
    }
}

//-----
//returns the sum of each this.species average fitness
getAvgFitnessSum() {
    var averageSum = 0;
    for (var s of this.species) {
        averageSum += s.averageFitness;
    }
    return averageSum;
}

//-----
//kill the bottom half of each this.species
cullSpecies() {
    for (var s of this.species) {
        s.cull(); //kill bottom half
        s.fitnessSharing(); //also while we're at it lets do fitness sharing
        s.setAverage(); //reset averages because they will have changed
    }
}

massExtinction() {
    for (var i = 5; i < this.species.length; i++) {
        // this.species.remove(i); //sad
        this.species.splice(i, 1);
        i--;
    }
}

//-----
//
// BATCH LEARNING
//-----
//update all the players which are alive
updateAliveInBatches() {
    let aliveCount = 0;
    for (var i = 0; i < this.players.length; i++) {
        if (this.playerInBatch(this.players[i])) {

            if (!this.players[i].dead) {
                aliveCount++;
            }
        }
    }
}

```

```

        this.players[i].look(); //get inputs for brain
        this.players[i].think(); //use outputs from neural network
        this.players[i].update(); //move the player according to the outputs from the neural network
        if (!showNothing && !showBest || i == 0) {
            this.players[i].show();
        }
        if (this.players[i].score > this.globalBestScore) {
            this.globalBestScore = this.players[i].score;
        }
    }
}

if (aliveCount == 0) {
    this.batchNo++;
}

}

playerInBatch(player) {
    for (var i = this.batchNo * this.worldsPerBatch; i < min((this.batchNo + 1) * this.worldsPerBatch, worlds.length); i++) {
        if (player.world == worlds[i]) {
            return true;
        }
    }

    return false;
}

stepWorldsInBatch() {
    for (var i = this.batchNo * this.worldsPerBatch; i < min((this.batchNo + 1) * this.worldsPerBatch, worlds.length); i++) {
        worlds[i].Step(1 / 30, 10, 10);
    }
}
//-----
//returns true if all the players in a batch are dead    sad
batchDead() {
    for (var i = this.batchNo * this.playersPerBatch; i < min((this.batchNo + 1) * this.playersPerBatch, this.players.length); i++) {
        if (!this.players[i].dead) {
            return false;
        }
    }
    return true;
}
}
}

```

[illegible]