

<i><b>Individual</b></i>
<ul style="list-style-type: none"><li>- <code>_chromosome</code>: <code>numpy.ndarray</code></li><li>- <code>_weights</code>: <code>numpy.ndarray</code></li><li>- <code>_fitness</code>: <code>float</code></li></ul>
<ul style="list-style-type: none"><li>+ <code>get_chromosome()</code>: <code>numpy.ndarray</code></li><li>+ <code>get_weights()</code>: <code>numpy.ndarray</code></li><li>+ <code>get_fitness()</code>: <code>float</code></li><li>+ <code>set_chromosome()</code>: <code>void</code></li><li>+ <code>set_weights()</code>: <code>void</code></li><li>+ <code>set_fitness()</code>: <code>void</code></li></ul>