Computer Assignment 3

CPE 261456 (Introduction to Computational Intelligence)


โดย

นายพีรณัฐ ธารทะเลทอง

รหัสนักศึกษา 550610530


เสนอ

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**ลักษณะการทำงานของระบบ**

เริ่มต้น เป็นการกำหนด จำนวนโครโมโซม , generation ในการ train และโครงสร้าง Neuron network ที่จะใช้ให้ GA จากนั้นจะทำการ initial โครโมโซม โดยการสุ่มค่าน้ำหนัก -1 ถึง 1 โดยโครโมโซมจะมีความยาวเท่ากับจำนวนน้ำหนักในโครงสร้าง Neuron network จากนั้นทำ 10% cross validation โดยในขั้นตอนการ train ในแต่ละ fold นั้น เริ่มจาก สุ่มเลือกโครโมโซม ทีละคู่เพื่อนำมา crossover โดยใช้การ crossover แบบหนึ่งจุดจากตรงกลางของโครโมโซม โดยได้ลูกจากการทำ crossover มา 90% จากทั้งหมด จากนั้นนำโครโมโซมลูกที่ได้จาการ crossover มารวมกับกลุ่มพ่อแม่และให้ชื่อว่ากลุ่ม p1 จากนั้นเลือกมาเป็น 60% ของ generation ถัดไป โดยเลือกโครโมโซม ที่มีค่า fitness ดีที่สุดจากทั้งหมด โดยค่า fitness นั้นหาจากจำนวนความถูกต้องของคำตอบจากข้อมูลใน training set และโครโมโซมที่เหลืออีก 40% ได้จากการสุ่มโครโมโซมในกลุ่ม p1 มาทำการ mutate โดยสุ่ม node ของ neuron network มา 30% จาก node ทั้งหมด และเพิ่มค่าน้ำหนักขาเข้า node นั้น โดยสุ่มจาก -1 ถึง 1 จากนั้นนำโครโมโซมที่มีค่า fitness ที่ดีที่สุดใน generation สุดท้ายมาทดสอบกับชุดข้อมูลทดสอบ เพื่อหาค่าความผิดพลาด เมื่อครบทุก fold ในขั้นตอน cross validation แล้วสุดท้ายจะได้ค่าความผิดพลาดเฉลี่ยที่บ่งบอกถึงสมรรถนะของรูปแบบ GA นี้

**ตัวอย่าง output ของโปรแกรม**

Neuron network 30-12-2

---------- Fold: 1 ----------

Generation 1 best accurate: 77.58284600389864%

Generation 2 best accurate: 77.58284600389864%

Generation 3 best accurate: 80.50682261208577%

Generation 4 best accurate: 88.10916179337232%

Generation 5 best accurate: 88.69395711500975%

Generation 6 best accurate: 88.69395711500975%

Generation 7 best accurate: 89.8635477582846%

Generation 8 best accurate: 89.8635477582846%

Generation 9 best accurate: 89.8635477582846%

Generation 10 best accurate: 89.8635477582846%

Generation 11 best accurate: 89.8635477582846%

Generation 12 best accurate: 91.2280701754386%

Generation 13 best accurate: 91.81286549707602%

Generation 14 best accurate: 92.00779727095517%

Generation 15 best accurate: 92.00779727095517%

Generation 16 best accurate: 92.20272904483431%

Generation 17 best accurate: 92.20272904483431%

Generation 18 best accurate: 92.20272904483431%

Generation 19 best accurate: 92.20272904483431%

Generation 20 best accurate: 92.20272904483431%

Generation 21 best accurate: 92.20272904483431%

Generation 22 best accurate: 92.20272904483431%

Generation 23 best accurate: 92.20272904483431%

Generation 24 best accurate: 92.5925925925926%

Generation 25 best accurate: 92.5925925925926%

Generation 26 best accurate: 92.5925925925926%

Generation 27 best accurate: 92.5925925925926%

Generation 28 best accurate: 92.5925925925926%

Generation 29 best accurate: 92.5925925925926%

Generation 30 best accurate: 92.5925925925926%

Generation 31 best accurate: 92.5925925925926%

Generation 32 best accurate: 92.5925925925926%

Generation 33 best accurate: 92.78752436647173%

Generation 34 best accurate: 92.78752436647173%

Generation 35 best accurate: 92.78752436647173%

Generation 36 best accurate: 93.17738791423002%

Generation 37 best accurate: 93.17738791423002%

Generation 38 best accurate: 93.17738791423002%

Generation 39 best accurate: 93.17738791423002%

Generation 40 best accurate: 93.17738791423002%

Generation 41 best accurate: 93.17738791423002%

Generation 42 best accurate: 93.17738791423002%

Generation 43 best accurate: 93.17738791423002%

Generation 44 best accurate: 93.17738791423002%

Generation 45 best accurate: 93.17738791423002%

Generation 46 best accurate: 93.17738791423002%

Generation 47 best accurate: 93.17738791423002%

Generation 48 best accurate: 93.17738791423002%

Generation 49 best accurate: 93.17738791423002%

Generation 50 best accurate: 93.17738791423002%

Testing accurate: 85.71428571428571%

---------- Fold: 2 ----------

Generation 1 best accurate: 86.93957115009746%

Generation 2 best accurate: 86.93957115009746%

Generation 3 best accurate: 87.91423001949317%

Generation 4 best accurate: 87.91423001949317%

Generation 5 best accurate: 87.91423001949317%

Generation 6 best accurate: 88.30409356725146%

Generation 7 best accurate: 88.88888888888889%

Generation 8 best accurate: 88.88888888888889%

Generation 9 best accurate: 88.88888888888889%

Generation 10 best accurate: 88.88888888888889%

Generation 11 best accurate: 89.8635477582846%

Generation 12 best accurate: 89.8635477582846%

Generation 13 best accurate: 90.05847953216374%

Generation 14 best accurate: 90.05847953216374%

Generation 15 best accurate: 90.44834307992203%

Generation 16 best accurate: 90.44834307992203%

Generation 17 best accurate: 90.44834307992203%

Generation 18 best accurate: 90.83820662768031%

Generation 19 best accurate: 90.83820662768031%

Generation 20 best accurate: 90.83820662768031%

Generation 21 best accurate: 90.83820662768031%

Generation 22 best accurate: 90.83820662768031%

Generation 23 best accurate: 90.83820662768031%

Generation 24 best accurate: 91.03313840155946%

Generation 25 best accurate: 91.2280701754386%

Generation 26 best accurate: 91.61793372319688%

Generation 27 best accurate: 91.61793372319688%

Generation 28 best accurate: 91.61793372319688%

Generation 29 best accurate: 91.61793372319688%

Generation 30 best accurate: 91.61793372319688%

Generation 31 best accurate: 91.61793372319688%

Generation 32 best accurate: 91.61793372319688%

Generation 33 best accurate: 91.61793372319688%

Generation 34 best accurate: 91.61793372319688%

Generation 35 best accurate: 91.61793372319688%

Generation 36 best accurate: 91.61793372319688%

Generation 37 best accurate: 91.61793372319688%

Generation 38 best accurate: 91.61793372319688%

Generation 39 best accurate: 91.61793372319688%

Generation 40 best accurate: 91.61793372319688%

Generation 41 best accurate: 91.61793372319688%

Generation 42 best accurate: 91.61793372319688%

Generation 43 best accurate: 92.00779727095517%

Generation 44 best accurate: 92.00779727095517%

Generation 45 best accurate: 92.00779727095517%

Generation 46 best accurate: 92.00779727095517%

Generation 47 best accurate: 92.00779727095517%

Generation 48 best accurate: 92.00779727095517%

Generation 49 best accurate: 92.00779727095517%

Generation 50 best accurate: 92.00779727095517%

Testing accurate: 92.85714285714286%

---------- Fold: 3 ----------

Generation 1 best accurate: 78.3625730994152%

Generation 2 best accurate: 80.70175438596492%

Generation 3 best accurate: 80.70175438596492%

Generation 4 best accurate: 80.70175438596492%

Generation 5 best accurate: 84.99025341130604%

Generation 6 best accurate: 88.10916179337232%

Generation 7 best accurate: 88.88888888888889%

Generation 8 best accurate: 88.88888888888889%

Generation 9 best accurate: 88.88888888888889%

Generation 10 best accurate: 89.66861598440546%

Generation 11 best accurate: 89.66861598440546%

Generation 12 best accurate: 89.66861598440546%

Generation 13 best accurate: 89.66861598440546%

Generation 14 best accurate: 89.66861598440546%

Generation 15 best accurate: 89.66861598440546%

Generation 16 best accurate: 89.66861598440546%

Generation 17 best accurate: 89.66861598440546%

Generation 18 best accurate: 90.25341130604288%

Generation 19 best accurate: 90.25341130604288%

Generation 20 best accurate: 90.64327485380117%

Generation 21 best accurate: 90.64327485380117%

Generation 22 best accurate: 91.42300194931774%

Generation 23 best accurate: 91.42300194931774%

Generation 24 best accurate: 91.42300194931774%

Generation 25 best accurate: 91.42300194931774%

Generation 26 best accurate: 91.42300194931774%

Generation 27 best accurate: 91.42300194931774%

Generation 28 best accurate: 91.42300194931774%

Generation 29 best accurate: 91.42300194931774%

Generation 30 best accurate: 91.42300194931774%

Generation 31 best accurate: 91.42300194931774%

Generation 32 best accurate: 91.61793372319688%

Generation 33 best accurate: 91.61793372319688%

Generation 34 best accurate: 91.61793372319688%

Generation 35 best accurate: 91.61793372319688%

Generation 36 best accurate: 92.00779727095517%

Generation 37 best accurate: 92.00779727095517%

Generation 38 best accurate: 92.00779727095517%

Generation 39 best accurate: 92.00779727095517%

Generation 40 best accurate: 92.00779727095517%

Generation 41 best accurate: 92.5925925925926%

Generation 42 best accurate: 92.5925925925926%

Generation 43 best accurate: 92.5925925925926%

Generation 44 best accurate: 92.5925925925926%

Generation 45 best accurate: 92.5925925925926%

Generation 46 best accurate: 92.5925925925926%

Generation 47 best accurate: 92.5925925925926%

Generation 48 best accurate: 92.5925925925926%

Generation 49 best accurate: 92.78752436647173%

Generation 50 best accurate: 92.98245614035088%

Testing accurate: 94.64285714285714%

---------- Fold: 4 ----------

Generation 1 best accurate: 78.94736842105263%

Generation 2 best accurate: 88.30409356725146%

Generation 3 best accurate: 90.44834307992203%

Generation 4 best accurate: 90.44834307992203%

Generation 5 best accurate: 90.44834307992203%

Generation 6 best accurate: 90.44834307992203%

Generation 7 best accurate: 90.44834307992203%

Generation 8 best accurate: 90.44834307992203%

Generation 9 best accurate: 90.83820662768031%

Generation 10 best accurate: 90.83820662768031%

Generation 11 best accurate: 90.83820662768031%

Generation 12 best accurate: 90.83820662768031%

Generation 13 best accurate: 90.83820662768031%

Generation 14 best accurate: 90.83820662768031%

Generation 15 best accurate: 90.83820662768031%

Generation 16 best accurate: 90.83820662768031%

Generation 17 best accurate: 90.83820662768031%

Generation 18 best accurate: 90.83820662768031%

Generation 19 best accurate: 90.83820662768031%

Generation 20 best accurate: 90.83820662768031%

Generation 21 best accurate: 90.83820662768031%

Generation 22 best accurate: 90.83820662768031%

Generation 23 best accurate: 90.83820662768031%

Generation 24 best accurate: 90.83820662768031%

Generation 25 best accurate: 90.83820662768031%

Generation 26 best accurate: 91.2280701754386%

Generation 27 best accurate: 91.2280701754386%

Generation 28 best accurate: 91.2280701754386%

Generation 29 best accurate: 91.2280701754386%

Generation 30 best accurate: 91.61793372319688%

Generation 31 best accurate: 91.61793372319688%

Generation 32 best accurate: 91.61793372319688%

Generation 33 best accurate: 92.78752436647173%

Generation 34 best accurate: 92.78752436647173%

Generation 35 best accurate: 92.78752436647173%

Generation 36 best accurate: 92.78752436647173%

Generation 37 best accurate: 92.78752436647173%

Generation 38 best accurate: 92.78752436647173%

Generation 39 best accurate: 92.78752436647173%

Generation 40 best accurate: 92.78752436647173%

Generation 41 best accurate: 92.78752436647173%

Generation 42 best accurate: 92.78752436647173%

Generation 43 best accurate: 92.78752436647173%

Generation 44 best accurate: 92.78752436647173%

Generation 45 best accurate: 92.78752436647173%

Generation 46 best accurate: 92.78752436647173%

Generation 47 best accurate: 92.78752436647173%

Generation 48 best accurate: 92.78752436647173%

Generation 49 best accurate: 92.98245614035088%

Generation 50 best accurate: 92.98245614035088%

Testing accurate: 92.85714285714286%

---------- Fold: 5 ----------

Generation 1 best accurate: 91.03313840155946%

Generation 2 best accurate: 91.03313840155946%

Generation 3 best accurate: 91.03313840155946%

Generation 4 best accurate: 91.03313840155946%

Generation 5 best accurate: 91.03313840155946%

Generation 6 best accurate: 91.03313840155946%

Generation 7 best accurate: 91.03313840155946%

Generation 8 best accurate: 91.03313840155946%

Generation 9 best accurate: 91.03313840155946%

Generation 10 best accurate: 91.03313840155946%

Generation 11 best accurate: 91.03313840155946%

Generation 12 best accurate: 91.03313840155946%

Generation 13 best accurate: 91.03313840155946%

Generation 14 best accurate: 91.03313840155946%

Generation 15 best accurate: 91.03313840155946%

Generation 16 best accurate: 91.03313840155946%

Generation 17 best accurate: 91.03313840155946%

Generation 18 best accurate: 91.03313840155946%

Generation 19 best accurate: 91.03313840155946%

Generation 20 best accurate: 91.2280701754386%

Generation 21 best accurate: 91.2280701754386%

Generation 22 best accurate: 91.2280701754386%

Generation 23 best accurate: 91.2280701754386%

Generation 24 best accurate: 91.2280701754386%

Generation 25 best accurate: 91.2280701754386%

Generation 26 best accurate: 91.2280701754386%

Generation 27 best accurate: 91.2280701754386%

Generation 28 best accurate: 91.2280701754386%

Generation 29 best accurate: 91.2280701754386%

Generation 30 best accurate: 91.2280701754386%

Generation 31 best accurate: 91.42300194931774%

Generation 32 best accurate: 91.42300194931774%

Generation 33 best accurate: 91.42300194931774%

Generation 34 best accurate: 91.42300194931774%

Generation 35 best accurate: 91.42300194931774%

Generation 36 best accurate: 91.42300194931774%

Generation 37 best accurate: 91.61793372319688%

Generation 38 best accurate: 91.61793372319688%

Generation 39 best accurate: 91.61793372319688%

Generation 40 best accurate: 91.61793372319688%

Generation 41 best accurate: 91.61793372319688%

Generation 42 best accurate: 91.61793372319688%

Generation 43 best accurate: 91.61793372319688%

Generation 44 best accurate: 91.61793372319688%

Generation 45 best accurate: 91.61793372319688%

Generation 46 best accurate: 91.61793372319688%

Generation 47 best accurate: 91.61793372319688%

Generation 48 best accurate: 91.61793372319688%

Generation 49 best accurate: 91.61793372319688%

Generation 50 best accurate: 91.61793372319688%

Testing accurate: 82.14285714285714%

---------- Fold: 6 ----------

Generation 1 best accurate: 77.77777777777777%

Generation 2 best accurate: 77.77777777777777%

Generation 3 best accurate: 81.48148148148148%

Generation 4 best accurate: 89.8635477582846%

Generation 5 best accurate: 90.25341130604288%

Generation 6 best accurate: 90.25341130604288%

Generation 7 best accurate: 90.25341130604288%

Generation 8 best accurate: 90.25341130604288%

Generation 9 best accurate: 92.20272904483431%

Generation 10 best accurate: 92.20272904483431%

Generation 11 best accurate: 92.20272904483431%

Generation 12 best accurate: 92.20272904483431%

Generation 13 best accurate: 92.20272904483431%

Generation 14 best accurate: 92.20272904483431%

Generation 15 best accurate: 92.20272904483431%

Generation 16 best accurate: 92.20272904483431%

Generation 17 best accurate: 92.20272904483431%

Generation 18 best accurate: 92.20272904483431%

Generation 19 best accurate: 92.20272904483431%

Generation 20 best accurate: 92.20272904483431%

Generation 21 best accurate: 92.20272904483431%

Generation 22 best accurate: 92.20272904483431%

Generation 23 best accurate: 92.5925925925926%

Generation 24 best accurate: 92.5925925925926%

Generation 25 best accurate: 92.5925925925926%

Generation 26 best accurate: 92.5925925925926%

Generation 27 best accurate: 92.5925925925926%

Generation 28 best accurate: 92.5925925925926%

Generation 29 best accurate: 92.5925925925926%

Generation 30 best accurate: 92.98245614035088%

Generation 31 best accurate: 92.98245614035088%

Generation 32 best accurate: 92.98245614035088%

Generation 33 best accurate: 92.98245614035088%

Generation 34 best accurate: 92.98245614035088%

Generation 35 best accurate: 92.98245614035088%

Generation 36 best accurate: 92.98245614035088%

Generation 37 best accurate: 92.98245614035088%

Generation 38 best accurate: 92.98245614035088%

Generation 39 best accurate: 92.98245614035088%

Generation 40 best accurate: 92.98245614035088%

Generation 41 best accurate: 92.98245614035088%

Generation 42 best accurate: 92.98245614035088%

Generation 43 best accurate: 92.98245614035088%

Generation 44 best accurate: 92.98245614035088%

Generation 45 best accurate: 92.98245614035088%

Generation 46 best accurate: 92.98245614035088%

Generation 47 best accurate: 92.98245614035088%

Generation 48 best accurate: 92.98245614035088%

Generation 49 best accurate: 92.98245614035088%

Generation 50 best accurate: 92.98245614035088%

Testing accurate: 87.5%

---------- Fold: 7 ----------

Generation 1 best accurate: 81.676413255360063%

Generation 2 best accurate: 81.67641325536063%

Generation 3 best accurate: 81.67641325536063%

Generation 4 best accurate: 82.06627680311891%

Generation 5 best accurate: 88.4990253411306%

Generation 6 best accurate: 88.4990253411306%

Generation 7 best accurate: 88.4990253411306%

Generation 8 best accurate: 90.44834307992203%

Generation 9 best accurate: 90.44834307992203%

Generation 10 best accurate: 90.44834307992203%

Generation 11 best accurate: 90.44834307992203%

Generation 12 best accurate: 90.44834307992203%

Generation 13 best accurate: 90.44834307992203%

Generation 14 best accurate: 90.44834307992203%

Generation 15 best accurate: 90.44834307992203%

Generation 16 best accurate: 90.44834307992203%

Generation 17 best accurate: 90.44834307992203%

Generation 18 best accurate: 90.44834307992203%

Generation 19 best accurate: 90.44834307992203%

Generation 20 best accurate: 90.44834307992203%

Generation 21 best accurate: 90.44834307992203%

Generation 22 best accurate: 90.44834307992203%

Generation 23 best accurate: 90.44834307992203%

Generation 24 best accurate: 90.44834307992203%

Generation 25 best accurate: 91.03313840155946%

Generation 26 best accurate: 91.2280701754386%

Generation 27 best accurate: 91.42300194931774%

Generation 28 best accurate: 91.42300194931774%

Generation 29 best accurate: 91.42300194931774%

Generation 30 best accurate: 91.42300194931774%

Generation 31 best accurate: 91.42300194931774%

Generation 32 best accurate: 91.61793372319688%

Generation 33 best accurate: 91.61793372319688%

Generation 34 best accurate: 91.61793372319688%

Generation 35 best accurate: 91.61793372319688%

Generation 36 best accurate: 91.61793372319688%

Generation 37 best accurate: 91.61793372319688%

Generation 38 best accurate: 91.61793372319688%

Generation 39 best accurate: 91.81286549707602%

Generation 40 best accurate: 91.81286549707602%

Generation 41 best accurate: 91.81286549707602%

Generation 42 best accurate: 91.81286549707602%

Generation 43 best accurate: 91.81286549707602%

Generation 44 best accurate: 91.81286549707602%

Generation 45 best accurate: 91.81286549707602%

Generation 46 best accurate: 91.81286549707602%

Generation 47 best accurate: 91.81286549707602%

Generation 48 best accurate: 91.81286549707602%

Generation 49 best accurate: 92.00779727095517%

Generation 50 best accurate: 92.00779727095517%

Testing accurate: 92.85714285714286%

---------- Fold: 8 ----------

Generation 1 best accurate: 83.23586744639377%

Generation 2 best accurate: 86.15984405458090%

Generation 3 best accurate: 86.15984405458090%

Generation 4 best accurate: 86.15984405458090%

Generation 5 best accurate: 86.15984405458090%

Generation 6 best accurate: 86.15984405458090%

Generation 7 best accurate: 91.03313840155946%

Generation 8 best accurate: 91.03313840155946%

Generation 9 best accurate: 91.03313840155946%

Generation 10 best accurate: 91.03313840155946%

Generation 11 best accurate: 91.03313840155946%

Generation 12 best accurate: 91.03313840155946%

Generation 13 best accurate: 91.03313840155946%

Generation 14 best accurate: 91.03313840155946%

Generation 15 best accurate: 91.03313840155946%

Generation 16 best accurate: 91.03313840155946%

Generation 17 best accurate: 91.03313840155946%

Generation 18 best accurate: 91.03313840155946%

Generation 19 best accurate: 91.03313840155946%

Generation 20 best accurate: 91.03313840155946%

Generation 21 best accurate: 91.03313840155946%

Generation 22 best accurate: 91.03313840155946%

Generation 23 best accurate: 91.03313840155946%

Generation 24 best accurate: 91.03313840155946%

Generation 25 best accurate: 91.2280701754386%

Generation 26 best accurate: 91.2280701754386%

Generation 27 best accurate: 91.2280701754386%

Generation 28 best accurate: 91.2280701754386%

Generation 29 best accurate: 91.2280701754386%

Generation 30 best accurate: 91.2280701754386%

Generation 31 best accurate: 91.2280701754386%

Generation 32 best accurate: 91.2280701754386%

Generation 33 best accurate: 91.2280701754386%

Generation 34 best accurate: 91.2280701754386%

Generation 35 best accurate: 91.2280701754386%

Generation 36 best accurate: 91.2280701754386%

Generation 37 best accurate: 91.2280701754386%

Generation 38 best accurate: 91.2280701754386%

Generation 39 best accurate: 91.2280701754386%

Generation 40 best accurate: 91.42300194931774%

Generation 41 best accurate: 91.42300194931774%

Generation 42 best accurate: 91.42300194931774%

Generation 43 best accurate: 91.42300194931774%

Generation 44 best accurate: 91.42300194931774%

Generation 45 best accurate: 91.61793372319688%

Generation 46 best accurate: 91.61793372319688%

Generation 47 best accurate: 91.61793372319688%

Generation 48 best accurate: 91.61793372319688%

Generation 49 best accurate: 91.61793372319688%

Generation 50 best accurate: 91.61793372319688%

Testing accurate: 92.85714285714286%

---------- Fold: 9 ----------

Generation 1 best accurate: 84.60038986354776%

Generation 2 best accurate: 86.1598440545809%

Generation 3 best accurate: 86.93957115009746%

Generation 4 best accurate: 88.69395711500975%

Generation 5 best accurate: 88.88888888888889%

Generation 6 best accurate: 88.88888888888889%

Generation 7 best accurate: 88.88888888888889%

Generation 8 best accurate: 90.25341130604288%

Generation 9 best accurate: 91.2280701754386%

Generation 10 best accurate: 91.2280701754386%

Generation 11 best accurate: 91.2280701754386%

Generation 12 best accurate: 91.2280701754386%

Generation 13 best accurate: 91.2280701754386%

Generation 14 best accurate: 91.2280701754386%

Generation 15 best accurate: 91.2280701754386%

Generation 16 best accurate: 91.2280701754386%

Generation 17 best accurate: 91.42300194931774%

Generation 18 best accurate: 91.42300194931774%

Generation 19 best accurate: 91.61793372319688%

Generation 20 best accurate: 91.61793372319688%

Generation 21 best accurate: 91.61793372319688%

Generation 22 best accurate: 91.61793372319688%

Generation 23 best accurate: 91.61793372319688%

Generation 24 best accurate: 91.61793372319688%

Generation 25 best accurate: 91.61793372319688%

Generation 26 best accurate: 91.61793372319688%

Generation 27 best accurate: 91.81286549707602%

Generation 28 best accurate: 92.39766081871345%

Generation 29 best accurate: 92.39766081871345%

Generation 30 best accurate: 92.39766081871345%

Generation 31 best accurate: 92.39766081871345%

Generation 32 best accurate: 92.39766081871345%

Generation 33 best accurate: 92.39766081871345%

Generation 34 best accurate: 92.5925925925926%

Generation 35 best accurate: 92.5925925925926%

Generation 36 best accurate: 92.5925925925926%

Generation 37 best accurate: 92.5925925925926%

Generation 38 best accurate: 92.5925925925926%

Generation 39 best accurate: 92.5925925925926%

Generation 40 best accurate: 92.5925925925926%

Generation 41 best accurate: 92.5925925925926%

Generation 42 best accurate: 92.5925925925926%

Generation 43 best accurate: 92.5925925925926%

Generation 44 best accurate: 92.5925925925926%

Generation 45 best accurate: 92.5925925925926%

Generation 46 best accurate: 92.5925925925926%

Generation 47 best accurate: 92.78752436647173%

Generation 48 best accurate: 92.78752436647173%

Generation 49 best accurate: 92.78752436647173%

Generation 50 best accurate: 92.78752436647173%

Testing accurate: 89.28571428571429%

---------- Fold: 10 ----------

Generation 1 best accurate: 86.15984405458091%

Generation 2 best accurate: 86.15984405458091%

Generation 3 best accurate: 86.54970760233918%

Generation 4 best accurate: 89.47368421052632%

Generation 5 best accurate: 89.47368421052632%

Generation 6 best accurate: 89.47368421052632%

Generation 7 best accurate: 89.47368421052632%

Generation 8 best accurate: 89.47368421052632%

Generation 9 best accurate: 89.47368421052632%

Generation 10 best accurate: 89.47368421052632%

Generation 11 best accurate: 89.47368421052632%

Generation 12 best accurate: 89.66861598440546%

Generation 13 best accurate: 91.03313840155946%

Generation 14 best accurate: 91.03313840155946%

Generation 15 best accurate: 91.03313840155946%

Generation 16 best accurate: 91.03313840155946%

Generation 17 best accurate: 92.20272904483431%

Generation 18 best accurate: 92.20272904483431%

Generation 19 best accurate: 92.20272904483431%

Generation 20 best accurate: 92.20272904483431%

Generation 21 best accurate: 92.20272904483431%

Generation 22 best accurate: 92.20272904483431%

Generation 23 best accurate: 92.20272904483431%

Generation 24 best accurate: 92.20272904483431%

Generation 25 best accurate: 92.20272904483431%

Generation 26 best accurate: 92.20272904483431%

Generation 27 best accurate: 92.20272904483431%

Generation 28 best accurate: 92.78752436647173%

Generation 29 best accurate: 92.78752436647173%

Generation 30 best accurate: 92.78752436647173%

Generation 31 best accurate: 92.78752436647173%

Generation 32 best accurate: 92.78752436647173%

Generation 33 best accurate: 92.78752436647173%

Generation 34 best accurate: 92.78752436647173%

Generation 35 best accurate: 92.78752436647173%

Generation 36 best accurate: 92.78752436647173%

Generation 37 best accurate: 92.78752436647173%

Generation 38 best accurate: 92.78752436647173%

Generation 39 best accurate: 92.78752436647173%

Generation 40 best accurate: 92.78752436647173%

Generation 41 best accurate: 92.78752436647173%

Generation 42 best accurate: 92.78752436647173%

Generation 43 best accurate: 92.78752436647173%

Generation 44 best accurate: 92.78752436647173%

Generation 45 best accurate: 92.78752436647173%

Generation 46 best accurate: 92.78752436647173%

Generation 47 best accurate: 92.98245614035088%

Generation 48 best accurate: 92.98245614035088%

Generation 49 best accurate: 92.98245614035088%

Generation 50 best accurate: 92.98245614035088%

Testing accurate: 92.85714285714286%

Error average: 9.642857142857142%

**การทดลอง**

1. ทำการทดลองโดยเปลี่ยน hidden layer ของ Neuron network เป็น 30-40-20-2, 30-15-7-2, 30-12-2, 30-10-5-2 โดยมีโครโมโซมใน GA เป็น 50 และ training สิ้นสุดใน generation ที่ 50 ได้ผลการทดลองดังนี้

   Neuron network 30-40-20-2

   ความผิดพลาดเฉลี่ยในการทำ 10% cross validation คือ 8.57142857142857%

   Neuron network 30-15-7-2

   ความผิดพลาดเฉลี่ยในการทำ 10% cross validation คือ 9.10714285714286%

   Neuron network 30-12-2

   ความผิดพลาดเฉลี่ยในการทำ 10% cross validation คือ 9.107142857142858%

   Neuron network 30-10-5-2

   ความผิดพลาดเฉลี่ยในการทำ 10% cross validation คือ 8.571428571428571%

2. ทำการทดลองโดยใช้ Neuron network 30-10-5-2 โดยการเพิ่มโครโมโซมเป็น 200 และเพิ่ม generation การ train เป็น 200 ได้ผลการทดลองดังนี้

   การเพิ่มโครโมโซมเป็น 200 ได้ความผิดพลาดเฉลี่ยคือ 9.464285714285714%

   การเพิ่ม generation การ train เป็น 200 ได้ความผิดพลาดเฉลี่ยคือ 8.928571428571427%

**วิเคราะห์ผลการทดลอง**

จากการทดลองการเปลี่ยน hidden layer ของ Neuron network จะเห็นได้ว่าความผิดพลาดเฉลี่ยนั้นมีค่าใกล้เคียงกันและโครงสร้างที่ดีที่สุดในการทดลองนี้คือ Neuron network 30-40-20-2 และในการทดลองการเพิ่มโครโมโซมและเพิ่ม generation การ train นั้น จากการทดลองแรก ใช้จำนวนโครโมโซมเป็น 50 และ generation การ train เป็น 50 ความผิดพลาดเฉลี่ยคือ 8.571428571428571% จะเห็นได้ว่าการเพิ่มจำนวนโครโมโซมและ generation การ train นั้นจะทำให้ความผิดพลาดเฉลี่ย เพิ่มขึ้น เพราะเป็นการเรียนรู้มากเกินไป (overtrain) ค่าความถูกต้องในขั้นตอน training จะมีค่ามากขึ้น แต่เมื่อนำข้อมูลทดสอบที่ต่างจากข้อมูล training มาทดสอบจะทำให้เกิดค่าความผิดพลาดมากกว่า

**Code** (https://github.com/porpeeranut/Computational_Intelligence_Assignment3 )

```
//          main.java

import java.io.BufferedReader;
import java.io.File;
import java.io.FileReader;
import java.io.IOException;
import java.util.ArrayList;
import java.util.Collections;


public class main {


        public static void main(String[] args) {
                String path = "..\\wdbc.data.txt";
                File file = new File(path);
                // feature 0-29, output 30
                ArrayList<Double[]> trainingSet = new ArrayList<Double[]>();
                try {
                        BufferedReader br = new BufferedReader(new FileReader(file));
                        String line;
                        while ((line = br.readLine()) != null) {
                                String[] data = line.split(",");
                                Double[] feature = new Double[31];
                                if (data[1].equals("M"))
                                        feature[30] = 0.0;
                                else
                                        feature[30] = 1.0;
                                for (int j = 2;j < data.length;j++) {
                                        feature[j-2] = Double.parseDouble(data[j]);
                                }
                                trainingSet.add(feature);
                        }
                        Collections.shuffle(trainingSet);
                        System.out.println(trainingSet.size());
                        br.close();
                } catch (IOException e) {
                        e.printStackTrace();
                }


                GA ga = new GA();
```

```
                //int[] MLP_struct = {30,40,20,2};
                //int[] MLP_struct = {30,15,7,2};
                //int[] MLP_struct = {30,12,2};
                int[] MLP_struct = {30,10,5,2};

                System.out.print("Neuron network ");
                for (int i = 0;i < MLP_struct.length;i++) {
                        if (i != 0)
                                System.out.print('-');
                        System.out.print(MLP_struct[i]);
                }
                ga.initChromosome(200, MLP_struct);
                double eav = 0;
                for (int c = 0;c < 10;c++) {
                        System.out.println("\n---------- Fold: "+(c+1)+" ----------");
                        int i = (int) (c*trainingSet.size()*0.1);
                        ArrayList<Double[]> test = new ArrayList<Double[]>(trainingSet.subList(i, (int)
(i+(trainingSet.size()*0.1))));
                        ArrayList<Double[]> train = (ArrayList<Double[]>) trainingSet.clone();
                        train.subList(i, (int) (i+(train.size()*0.1))).clear();
                        ga.train(50, (ArrayList<Double[]>) train);
                        eav += ga.test(test);
                }
                System.out.println("Error average: "+eav/10+"%");
        }
}
```

```
//          GA.java

import java.util.ArrayList;
import java.util.Collections;
import java.util.Comparator;
import java.util.LinkedHashSet;
import java.util.Random;
import java.util.Set;

public class GA {

        ArrayList<Chromosome> initChromosome = new ArrayList<Chromosome>();
        ArrayList<Chromosome> chromosomeList = new ArrayList<Chromosome>();
        ArrayList<Double[]> trainingSet;
        int[] MLP_struct;
        double wMin = -1.0;
        double wMax = 1.0;

        public void initChromosome(int amount, int[] MLP_struct){
                this.MLP_struct = MLP_struct;
                int chromoLen = 0;
                for (int i = 1;i < MLP_struct.length;i++) {
                        chromoLen += MLP_struct[i-1]*MLP_struct[i];
                }
                for (int a = 0;a < amount;a++) {
                        Double[] gene = new Double[chromoLen];
                        for (int g = 0;g < gene.length;g++) {
                                gene[g] = wMin + (wMax - wMin)*new Random().nextDouble();
                        }
                        initChromosome.add(new Chromosome(gene));
                }
        }

        public void train(int maxGeneration, ArrayList<Double[]> trainingSet){
                this.trainingSet = trainingSet;
                chromosomeList = (ArrayList<Chromosome>) initChromosome.clone();
                computeFitnessInList(chromosomeList);
                for (int g = 0;g < maxGeneration;g++) {
                        ArrayList<Chromosome> selected = randomSelect((int) (chromosomeList.size()*0.9));
                        ArrayList<Chromosome> crossed = crossover(selected, (int) (selected.size()*0.9));
```

```
                ArrayList<Chromosome> pool = (ArrayList<Chromosome>) chromosomeList.clone();
                pool.addAll(crossed);
                //ArrayList<Chromosome> mutated = mutate(pool, (int) (pool.size()*0.7));
                int mutateAmount = (int) (chromosomeList.size()*0.4);
                ArrayList<Chromosome> mutated = mutate(pool, mutateAmount);
                //pool.addAll(mutated);
                Collections.sort(pool, new Comparator<Chromosome>() {
            public int compare(Chromosome c1, Chromosome c2) {
                return (int) (c2.fitness - c1.fitness);
            }
        });
                chromosomeList = new ArrayList<Chromosome>(pool.subList(0, chromosomeList.size()-
mutateAmount));
                //chromosomeList = new ArrayList<Chromosome>(pool.subList(0, chromosomeList.size()));
                chromosomeList.addAll(mutated);
                System.out.println("Generation "+(g+1)+" best accurate:
"+(chromosomeList.get(0).fitness*100/trainingSet.size())+"%");
            }

            Collections.sort(chromosomeList, new Comparator<Chromosome>() {
        public int compare(Chromosome c1, Chromosome c2) {
            return (int) (c2.fitness - c1.fitness);
        }
    });
            for (Chromosome chromosome : chromosomeList) {
                //System.out.println(chromosome.fitness+" "+(chromosome.fitness*100/trainingSet.size()));
            }
    }

    public double test(ArrayList<Double[]> testSet){
            MLP mlp = new MLP(MLP_struct);
            Chromosome bestChomos = chromosomeList.get(0);
            mlp.setWeightFromChromosome(bestChomos);
            bestChomos.fitness = 0.0;
            for (Double[] data : testSet) {
                if (mlp.computeForward(data))
                    bestChomos.fitness += 1;
            }
            //computeFitness(chromosomeList.get(0));
            System.out.println("Testing accurate: "+(bestChomos.fitness*100/testSet.size())+"%");
```

```
		return 100-(bestChomos.fitness*100/testSet.size());
	}

	private void computeFitnessInList(ArrayList<Chromosome> chromosomeList) {
		for (Chromosome chromosome : chromosomeList) {
			computeFitness(chromosome);
		}
	}

	private void computeFitness(Chromosome chromosome) {
		MLP mlp = new MLP(MLP_struct);
		mlp.setWeightFromChromosome(chromosome);
		chromosome.fitness = 0.0;
		for (Double[] data : trainingSet) {
			if (mlp.computeForward(data))
				chromosome.fitness += 1;
		}
	}

	private ArrayList<Chromosome> randomSelect(int amount) {
		ArrayList<Chromosome> selected = new ArrayList<Chromosome>();
		for (int a = 0;a < amount;a++) {
			int i = new Random().nextInt(chromosomeList.size());
			selected.add(chromosomeList.get(i));
		}
		return selected;
	}

	private ArrayList<Chromosome> crossover(ArrayList<Chromosome> selected, int amount) {
		ArrayList<Chromosome> crossed = new ArrayList<Chromosome>();
		for (int a = 0;a < amount;a++) {
			int i1 = new Random().nextInt(selected.size());
			int i2 = new Random().nextInt(selected.size());
			Chromosome daddy = selected.get(i1);
			Chromosome mommy = selected.get(i2);
			int chromosomeLen = selected.get(0).gene.length;
			int helfLen = chromosomeLen/2;
			Double[] gene = new Double[chromosomeLen];
			for (int i = 0;i < helfLen;i++) {
				gene[i] = daddy.gene[i];
```

```
                }
                for (int i = helfLen;i < chromosomeLen;i++) {
                        gene[i] = mommy.gene[i];
                }
                crossed.add(new Chromosome(gene));
        }
        computeFitnessInList(crossed);
        return crossed;
    }

    private ArrayList<Chromosome> mutate(ArrayList<Chromosome> pool, int amount) {
        ArrayList<Chromosome> mutated = new ArrayList<Chromosome>();
        Random rng = new Random();
        Set<Integer> index = new LinkedHashSet<Integer>();
        while (index.size() < amount)
        {
                //      random index
           Integer next = rng.nextInt(pool.size());
           index.add(next);
        }
        int k = 0;
        for (int i : index) {
                Double[] gene = pool.get(i).gene.clone();
                int mutateRate = (int) (pool.get(i).gene.length * 0.3);
                for (int m = 0;m < mutateRate;m++) {
                        int w = rng.nextInt(pool.get(i).gene.length);
                        gene[w] += wMin + (wMax - wMin)*new Random().nextDouble();
                        if (gene[w] > wMax)
                                gene[w] = wMax;
                        if (gene[w] < wMin)
                                gene[w] = wMin;
                }
                mutated.add(new Chromosome(gene));
                computeFitness(mutated.get(k++));
        }
        return mutated;
    }
}
```

```
//          MLP.java

public class MLP {

        int[] MLP_struct;
        double[][][] weight;  // layer, maxnode, maxnode
        int maxNode = 0;

        public MLP(int[] MLP_struct) {
                this.MLP_struct = MLP_struct;
                int allNode = 0;
                for (int struc : MLP_struct) {
                        if (maxNode < struc)
                                maxNode = struc;
                        allNode += struc;
                }
                weight = new double[MLP_struct.length-1][maxNode][maxNode];
        }

        public void setWeightFromChromosome(Chromosome chromosome) {
                int g = 0;
                for (int l = 0;l < MLP_struct.length-1;l++) {
                        for (int n1 = 0;n1 < MLP_struct[l];n1++) {
                                for (int n2 = 0;n2 < MLP_struct[l+1];n2++) {
                                        weight[l][n1][n2] = chromosome.gene[g];
                                        g++;
                                }
                        }
                }
        }

        public boolean computeForward(Double[] data) {
                Double[][] y = new Double[MLP_struct.length][maxNode];    // layer, node
                y[0] = data;
                for (int L = 1;L < MLP_struct.length;L++) {
                        for (int n2 = 0;n2 < MLP_struct[L];n2++) {
                                Double v = 0.0;
                                for (int n1 = 0;n1 < MLP_struct[L-1];n1++) {
                                        v += y[L-1][n1] * weight[L-1][n1][n2];
                                }
```

```
                        //y[L][n2] = sigmoid(v);
                        y[L][n2] = Math.tanh(v);
                    }
                }
                if (data[30] == 1.0) {
                    //          "B" 0 1
                    if (y[MLP_struct.length-1][0] < y[MLP_struct.length-1][1])
                        return true;
                    else
                        return false;
                } else {
                    //          "M" 1 0
                    if (y[MLP_struct.length-1][0] > y[MLP_struct.length-1][1])
                        return true;
                    else
                        return false;
                }
            }

            private Double sigmoid(Double v) {
                return 1.0 / (1.0 + Math.exp(-v));
            }
}
```

```
//          Chromosome.java

public class Chromosome {

            public Double[] gene;
            public Double fitness = 0.0;

            public Chromosome(Double[] gene) {
                this.gene = gene;
            }
}
```