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Chromosome size – 42;

Capacity – 100;

Epoch – 50;

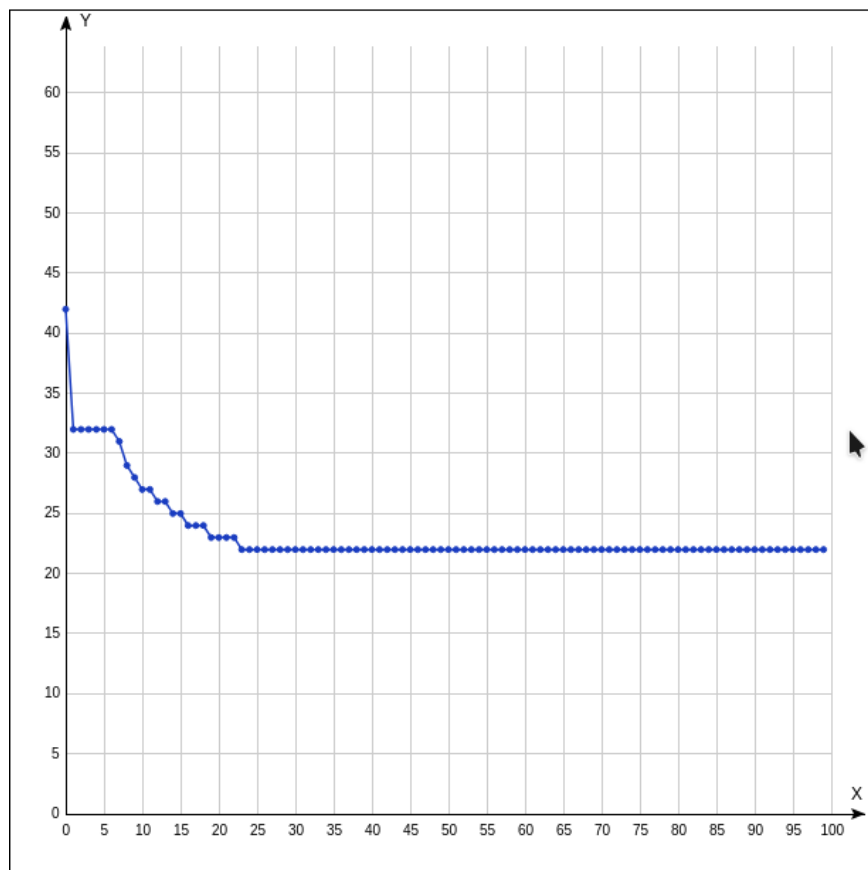
with mutation;

Onepoint crossover;

Truncle selection;

X generations

Y — error



Best network (start): 0.5456212 0.44264054 0.072345614 0.968213 0.97299993 0.4165203
0.9419711 0.46587026 0.07849419 0.4333101 0.84688497 0.56225383 0.3317436 0.1033566
0.40506822 0.3687924 0.34950835 0.7360943 0.55638725 0.88603693 0.7861397 0.16891843
0.6794174 0.09630722 0.29712468 0.39154583 0.33142322 0.7592533 0.74782705 0.27072102
0.687652 0.36509848 0.3789326 0.7731133 0.5762136 0.20631003 0.6374505 1.5205145E-4
0.08940798 0.50501186 0.8762369 0.12710059

Best network (10 generation): 0.7388291 0.99964464 0.39091504 0.81108886 0.76174325
0.62005556 0.36042124 0.9557455 0.24452579 0.120577395 0.3569411 0.40869498 0.95374
0.80706245 0.20638496 0.3860618 0.28198254 0.25567883 0.31531715 0.026140273 0.16370076
0.18330824 0.61856675 0.48481148 0.6684862 0.014447629 0.41877925 0.46704674 0.88221323
0.39763862 0.74882126 0.34414953 0.60882926 0.72660077 0.66762304 0.8668563 0.26870656
0.31269622 0.1676572 0.9886519 0.69589823 0.44865912

Best network (end): 0.6051165 0.30464655 0.6720695 0.52645606 0.5675815 0.41825616
0.29906678 0.7633289 0.18797165 0.21064585 0.4437796 0.97885936 0.38925958 0.69498295
0.38548917 0.4930178 0.91597193 0.379391 0.6325027 0.081130445 0.48945755 0.8950965
0.24120843 0.03933513 0.76550794 0.1013214 0.3634078 0.48738718 0.7127568 0.36924475
0.21097589 0.4952708 0.7955454 0.3826558 0.9362069 0.105920136 0.8142917 0.12799132
0.57276934 0.06466651 0.31101573 0.9541879

Conclusion: Algorithm can achieve 66% accuracy