

Evolving Neural Networks for Statistical Decision Theory

Michal Vaľko¹ Mgr. Radoslav Harman PhD.²

¹Department of Applied Informatics
Faculty of Mathematics, Physics and Informatics
Comenius University

²Department of Applied Mathematics and Statistics
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Master Thesis Defense, 2005



Outline

- 1 Introduction
 - Master Thesis Goals
- 2 Methods
 - JASTAP — Biologically Plausible NN Model
 - Inter-spike Intervals
 - Decisioning With NNs
 - Evolution
- 3 Decision Problems
 - More Frequent Input
 - Hypothesis Testing of Frequency
 - More Regular Input
 - Hypothesis Testing of Regularity



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Master Thesis Goals

- 1 Explore statistical decisioning in NNs
- 2 Analyze the abilities of simple network structures
- 3 Try to evolve the networks useful for statistical decisioning of mean rates and regularities
- 4 Methods: JASTAP and GA's



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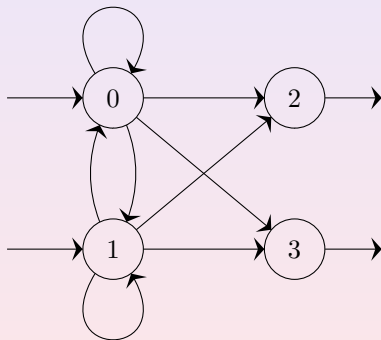


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What is JASTAP?

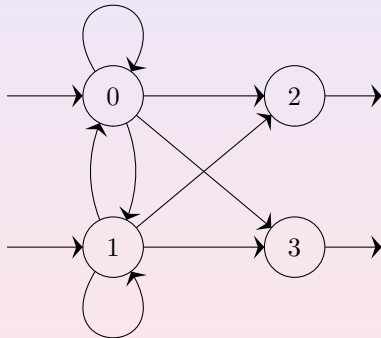


Characteristics

- spiking neuron model
- respects physiological aspects of a real neuron
- weights, thresholds, latencies, PSP's, firing rates



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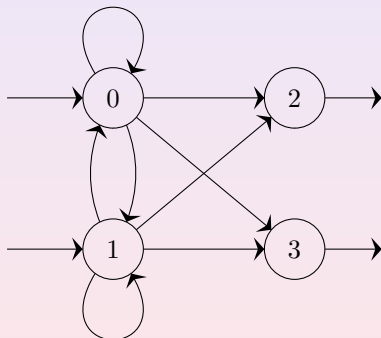


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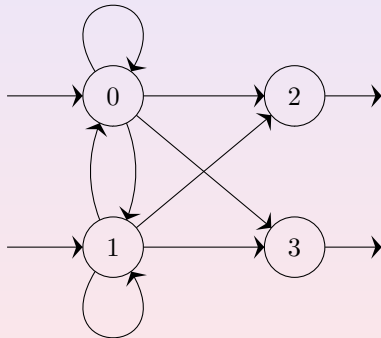


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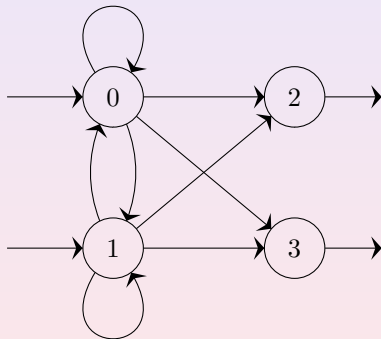


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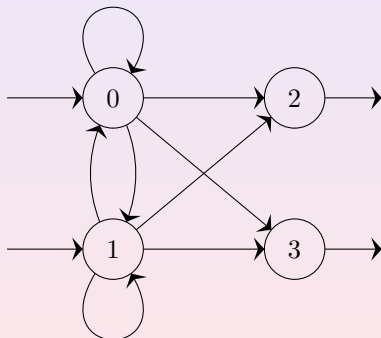


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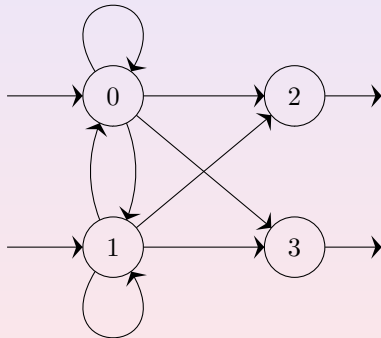


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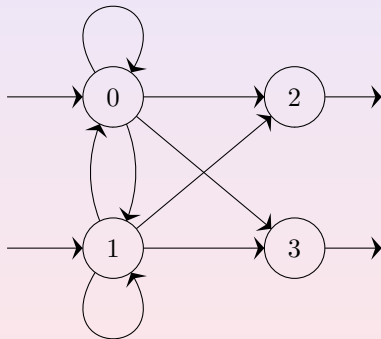


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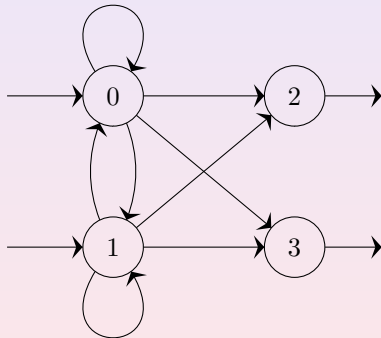


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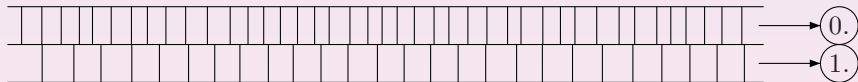
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Information Processing

Encoding

- JASTAP works with **temporal code**
- information is coded in inter-spike intervals



PSP Definition

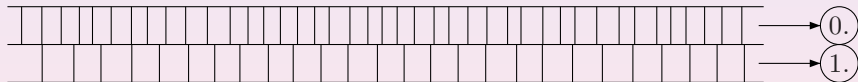
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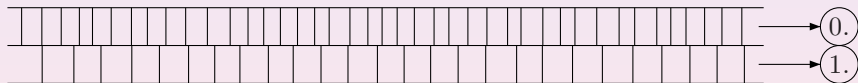
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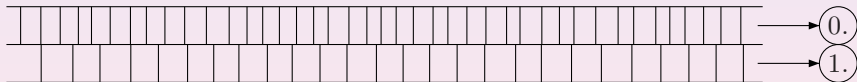
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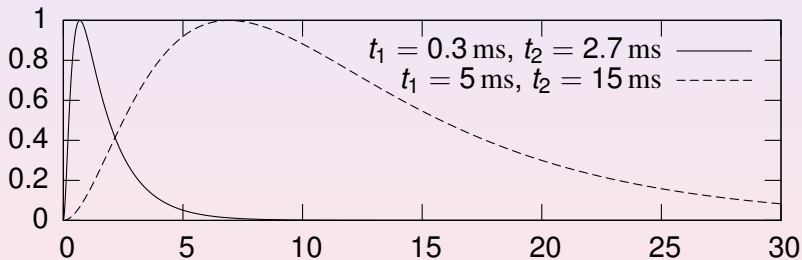


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Postsynaptic Potential



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Gamma Distribution

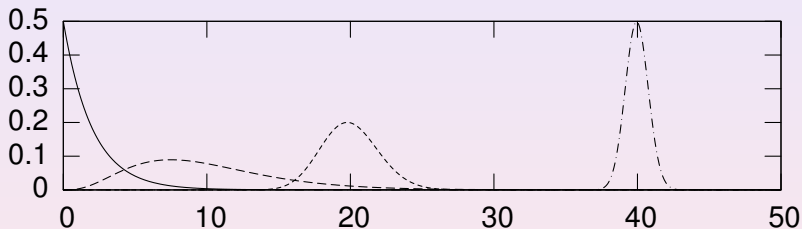
Definition

A random variable Z has the **Gamma distribution**, if the probabilistic density function of Z is

$$f(z) = \frac{z^{\alpha-1} e^{-\frac{z}{\beta}}}{\beta^{\alpha} \Gamma(\alpha)} \quad \alpha, \beta > 0, z \geq 0, \text{ and we denote it as } \mathcal{G}(\alpha, \beta)$$



Gamma Distribution



$c_V = 1, \overline{isi} = 2 \text{ ms}$ ———
 $c_V = 0.5, \overline{isi} = 10 \text{ ms}$ - - - - -
 $c_V = 0.1, \overline{isi} = 20 \text{ ms}$ ·····
 $c_V = 0.02, \overline{isi} = 40 \text{ ms}$ - · - · -



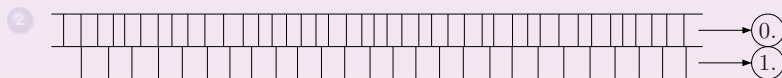
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How it works?

- 1 input temporal code is generated from random Gamma distribution draws

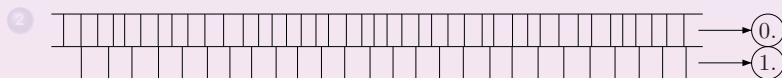


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- 4 if any of the output neurons fires, it is taken as a decision



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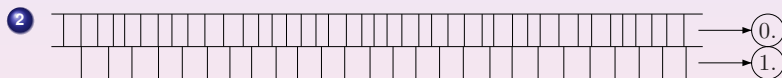


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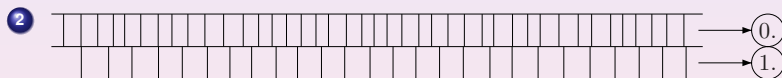


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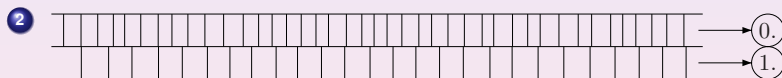


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Evolution Set Up

What to evolve?

weights absolutely

latencies important for time-related patterns

PSP shapes expands the search space, slower decay chosen

thresholds no comment

fire rates not evolved, $t_{\min} := 1 \text{ ms}$, $t_{\max} := 10 \text{ ms}$



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Genetic Algorithm

- Gray binary coding
- recombination: multipoint crossover
- mutation: p -scaled
- 2 phases: second is for *fine-tuning* from the *seed*
- **fitness function** → crucial issue



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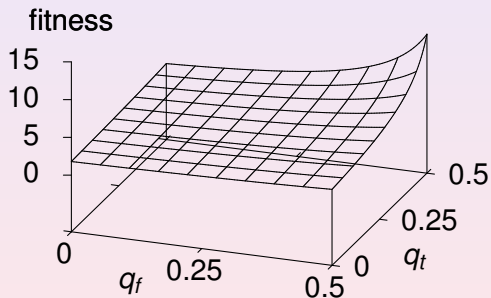
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Fitness Functions: Overall Ratio



overall ratio

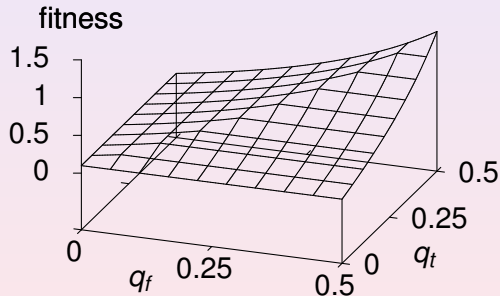
$$1/(1 + \varepsilon - q_t - q_f)$$



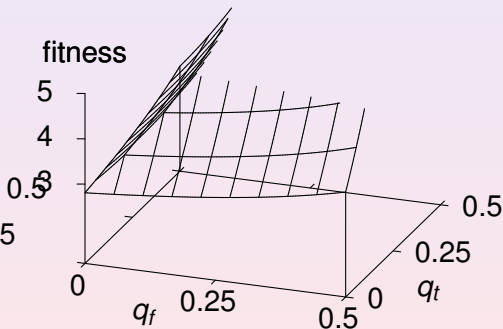
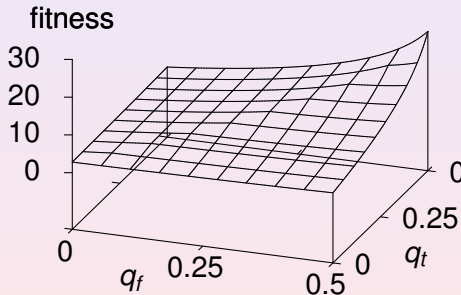
Fitness Functions: One Side Minimum

one side minimum

$$1/(1 - \min(q_t, q_f)) - 1$$



Fitness Functions: Combined



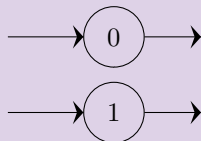
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Theoretical Strategies: Description

copy machine



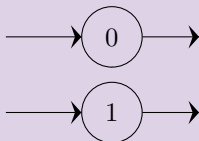
event counting

it makes the decision that the more frequent input is the one with the more events observed



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Theoretical Strategies: Defining the Bounds

copy machine

lower \bar{isi}	higher \bar{isi}	ratio
30 ms	40 ms	57.14 %
20 ms	40 ms	66.67 %
10 ms	40 ms	80.00 %
20 ms	30 ms	60.00 %
10 ms	30 ms	75.00 %
10 ms	20 ms	66.67 %

event counting

lower \bar{isi}	higher \bar{isi}	ratio
30 ms	40 ms	72.45 %
20 ms	40 ms	94.51 %
10 ms	40 ms	99.99 %
20 ms	30 ms	83.97 %
10 ms	30 ms	99.91 %
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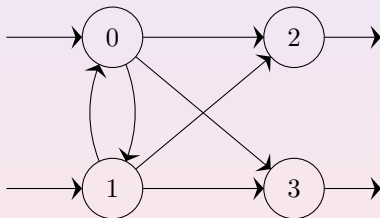
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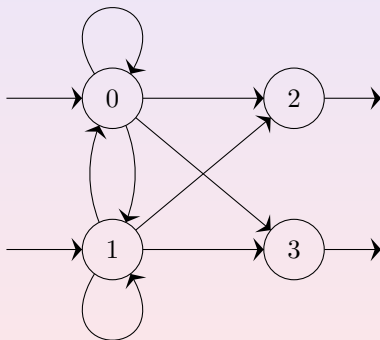
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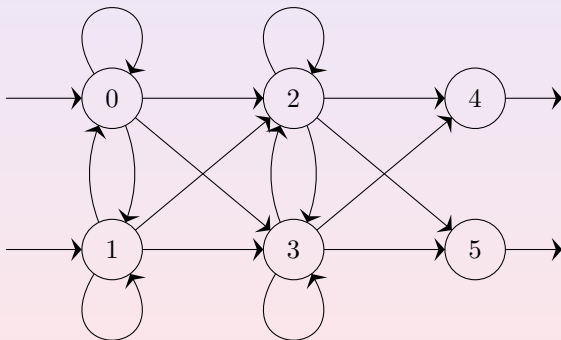
Network Structures: A



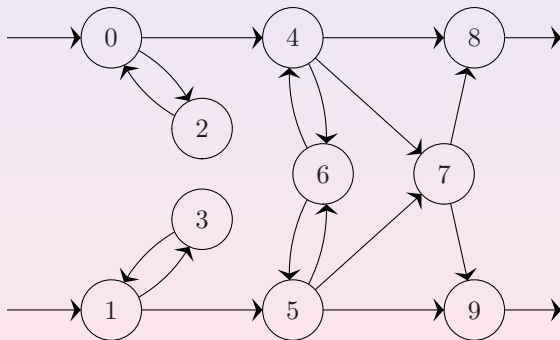
Network Structures: B



Network Structures: C



Network Structures: D

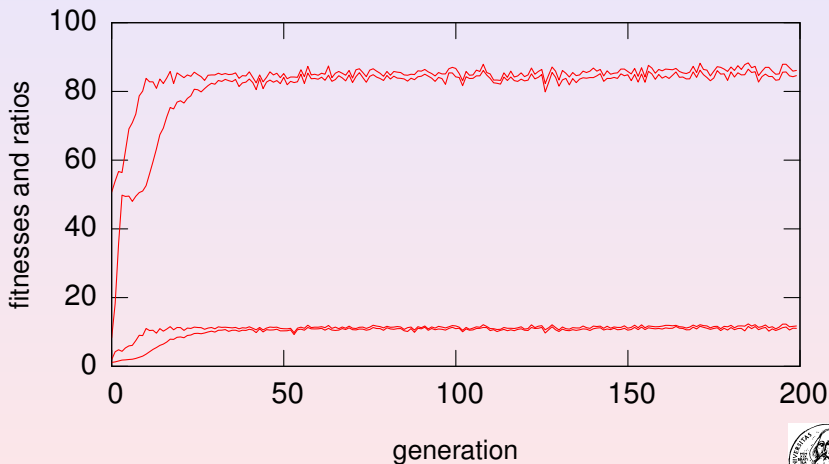


Network Structures: Comparison

low \overline{isi}	high \overline{isi}	copy	A	B	C	D	event
30 ms	40 ms	57.14	60.12	62.18	59.03	61.31	72.45
20 ms	40 ms	66.67	88.11	87.56	87.67	81.32	94.51
10 ms	40 ms	80.00	99.81	99.65	99.25	99.58	99.99
20 ms	30 ms	60.00	66.92	71.52	69.65	67.27	83.97
10 ms	30 ms	75.00	99.35	99.19	99.14	98.08	99.91
10 ms	20 ms	66.67	95.04	92.74	94.12	91.75	98.83
$\langle 10 \text{ ms}, 40 \text{ ms} \rangle$		60.22	66.66	66.68	68.48	65.75	79.68



Evolution curves: \overline{isi} 10 vs. 20 ms, $c_v = 1$

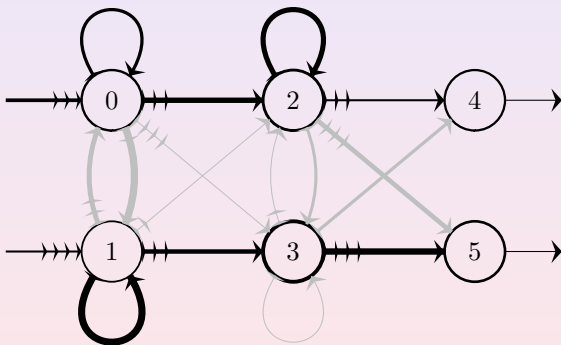


Comparison: Different c_v s (in %)

low \overline{isi}	high \overline{isi}	0.02	0.1	0.5	1	$\langle 0, 1 \rangle$
30 ms	40 ms	100.00	100.00	77.17	63.26	73.29
20 ms	40 ms	100.00	100.00	98.07	87.67	95.61
10 ms	40 ms	100.00	100.00	99.99	99.69	99.85
20 ms	30 ms	100.00	99.96	90.78	70.94	87.62
10 ms	30 ms	100.00	100.00	99.94	99.14	99.77
10 ms	20 ms	100.00	100.00	99.85	94.25	98.60
$\langle 10 \text{ ms}, 40 \text{ ms} \rangle$		98.02	92.51	78.15	72.00	66.01



Example: evolved network for \overline{isi} : 20 vs. 30 ms, $c_v = 1$



Results

Strategies

- information paths
- competing
- redundancy handling
- copy machine principle
- activity routing



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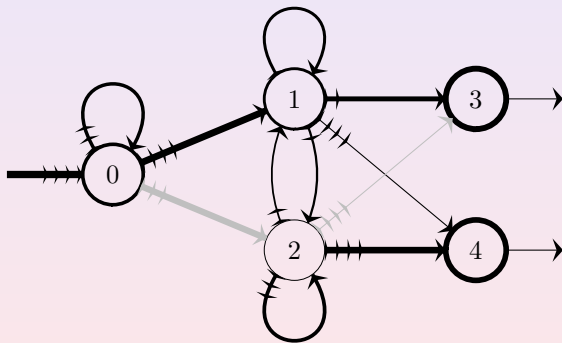


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Network Structure for Hypothesis Testing

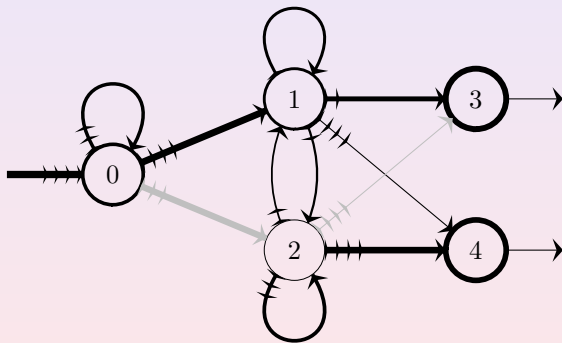


Strategies

- gap detection
- perfect timing



Network Structure for Hypothesis Testing



Strategies

- gap detection
- perfect timing



Results: $\overline{isi} \in_R \langle 10 \text{ ms}, 40 \text{ ms} \rangle, 300 \text{ ms}$

H_0	$c_V = 0.02$	$c_V = 0.1$	$c_V = 0.5$	$c_V = 1$	$\langle 0, 1 \rangle$
$\overline{isi} < 20 \text{ ms}$	98.57 %	94.72 %	89.48 %	57.20 %	73.91 %
$\overline{isi} < 25 \text{ ms}$	99.00 %	95.04 %	85.59 %	60.66 %	73.78 %
$\overline{isi} < 30 \text{ ms}$	98.97 %	94.64 %	67.53 %	56.86 %	74.21 %



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Results

low	high	10 ms	20 ms	30 ms	40 ms	$\langle 10, 40 \rangle$
0.02	0.50	99.93 %	99.75 %	98.90 %	98.74 %	74.52 %
0.02	1.00	99.95 %	99.61 %	97.99 %	98.32 %	88.85 %
0.50	1.00	83.05 %	76.55 %	77.44 %	64.32 %	67.08 %
$\langle 0.00, 1.00 \rangle$		70.99 %	67.85 %	67.78 %	63.96 %	55.07 %

Strategies

- close events
- distant events



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Outline

- 1 Introduction
 - Master Thesis Goals
- 2 Methods
 - JASTAP — Biologically Plausible NN Model
 - Inter-spike Intervals
 - Decisioning With NNs
 - Evolution
- 3 **Decision Problems**
 - More Frequent Input
 - Hypothesis Testing of Frequency
 - More Regular Input
 - **Hypothesis Testing of Regularity**



Results

H_0	10 ms	20 ms	30 ms	40 ms	$\langle 10, 40 \rangle$
$c_v < 0.5$	83.04 %	76.28 %	76.20 %	74.99 %	66.31 %

One Neuron Strategies

- memory in latency
- from regularity to frequency
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Conclusions

- JASTAP is able to model decision making.
- We have found decision makers for comparing and statistical testing of mean and c_v of Gamma distributions
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Further work

- Enhance the evolution.
- Evolve the topologies.
- Evolve *PSPs* and firing rates.
- Speed preferences, faster decision makers.



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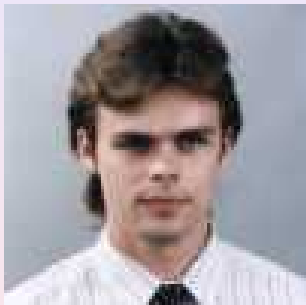
Obsah dodatku

4

Dodatok

- Školiteľský posudok
- Oponentský posudok
- Odovede na otázky oponenta
- Voľná diskusia





Mgr. Radoslav Harman, PhD.

Department of Applied
Mathematics and Statistics
Faculty of Mathematics,
Physics and Informatics
Comenius University





Ing. Igor Farkaš, PhD.

Department of Applied Informatics
Faculty of Mathematics,
Physics and Informatics
Comenius University



Oponent'ská otázka č. 1

Otázka

Autor si zvolil model JASTAP, no patrilo by sa aspoň v referenciách spomenúť, že existuje celá škála iných, etablovaných, biologicky prijateľných modelov neurónu (pozri napr. prehľadový článok Izhikevich E., IEEE Trans. on Neural Networks, 15(5), 2004). Okrem toho, čo znamená tá skratka?

Odpoveď

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- Jančo, Stavrovský, Pavlásek
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Niektoré symboly neboli vysvetlené, napr. str. 8: predpokladám, že $k=1$; pracuje sa v princípe v modeli aj s inou hodnotou k ? Str. 10: Čo je $\Gamma(a)$ pri gamma distribúcii $f(z)$?

Odpoveď

- k je normovacia konštanta, je hodnota je $1/(\text{maximálna hodnota PSP})$
- $\Gamma(z) = \int_0^{\infty} t^{z-1} e^{-t} dt$
- $\Gamma(n+1) = n\Gamma(n) = \dots = n! \Gamma(1) = n!$
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Otázka

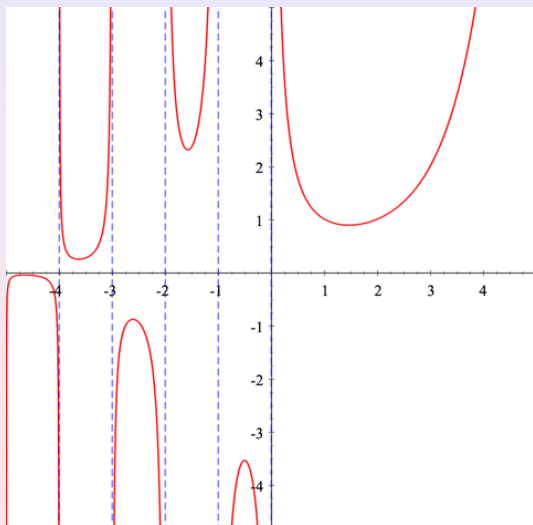
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Oponentská otázka č. 2



Oponentská otázka č. 3

Otázka

Rozdiely medzi jednotlivými modelmi AD (tab.4.3, 4.4) vyzerať by mali byť minimálne. Otázka je, či sú štatisticky významné. Pomohli by tu štatistické testy?

Odpoveď

Pozrime si znova tabuľku...



Oponentská otázka č. 3

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Odpoveď

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Oponentská otázka č. 3

vyššie \overline{isi}	nižšie \overline{isi}	A	B	C	D	rozdiel
30 ms	40 ms	60.12	62.18	59.03	61.31	2.06
20 ms	40 ms	88.11	87.56	87.67	81.32	6.79
10 ms	40 ms	99.81	99.65	99.25	99.58	0.40
20 ms	30 ms	66.92	71.52	69.65	67.27	4.60
10 ms	30 ms	99.35	99.19	99.14	98.08	1.27
10 ms	20 ms	95.04	92.74	94.12	91.75	3.29
$\langle 10 \text{ ms}, 40 \text{ ms} \rangle$		66.66	66.68	68.48	65.75	2.73



Oponent'ská otázka č. 3

Otázka

Rozdiely medzi jednotlivými modelmi AD (tab. 4.3, 4.4) vyzerajú byť minimálne. Otázka je, či sú štatisticky signifikantné. Pomohli by tu štatistické testy?

Odpoveď

Je pravdou, že testovanie jedincov kvôli ohodnoteniu počas simulácií bolo iba 50 testami z rýchlostostných dôvodov. V tabuľke sú však uvedené výsledky vypočítané z 10 000 testov a chyby sú na úrovni stotín.



Oponentská otázka č. 4

Otázka

Obr. 4.3: krivka pre *avg* kopíruje tú pre *best*. Očakával by som, že ako priemer bude *avg* hladká.

Odpoveď

Pozrime si dotýčný graf. . .



Oponentská otázka č. 4

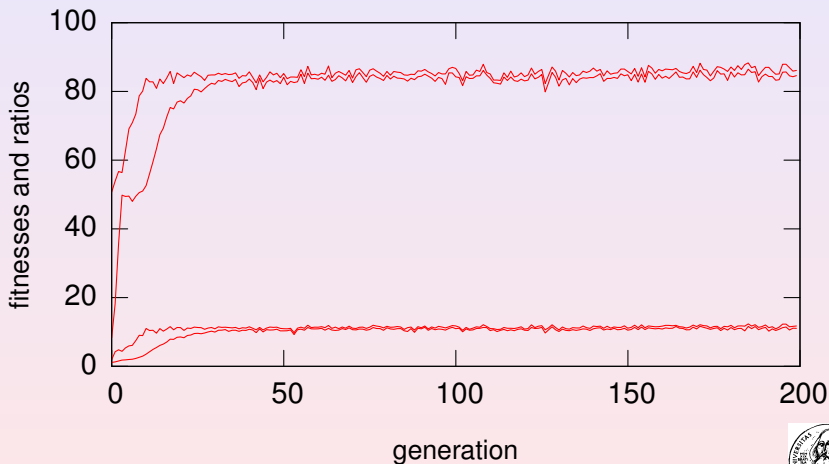
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Odpoveď

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Evolution curves: \overline{isi} 10 vs. 20 ms, $c_v = 1$ 

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Odpoveď

- skoky v grafe súvisia z generovaním úplne novej trénovacej sady pre každú generáciu
- na **ne**hladkosť má vplyv aj elitizmus
- čiarkovaný priebeh v tlačenej verzii DP



Oponentská otázka č. 5

Otázka

Z textu som nedokázal vydedukovať, čo znamenajú tie impulzy (prečo 4 línie), napr. obr. 4.4.

Odpoveď

Línie nad neurónmi znamenajú vstupy zo synáps. Štyri sú preto lebo zobrazované štruktúry majú štyri vstupy. Ak je neurón zároveň vstupným, prvá *línia* znázorňuje vonkajší vstup.



Oponentská otázka č. 5

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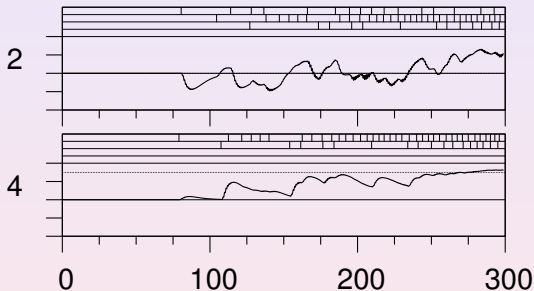
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Oponentská otázka č. 5



Obrázok: gap detection — a yes decision



Oponentská otázka č. 6

Otázka

Autor rieši biologicky relevantný problém biologicky relevantnými prostriedkami, avšak použil "fylogenetický" prístup (GA) na riešenie "ontologického" problému (učenie). Je to odôvodniteľné problémom návrhu vhodného tradičného algoritmu učenia (napr. na báze Hebbovho učenia), hoci tento prístup by bol zrejme viac biologicky prijateľný ako GA.

Odpoveď

Je to presne tak, ale cieľom bolo skôr nájsť štruktúry schopné rozhodovania než zistiť ako takéto štruktúry vznikli.



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Voľná diskusia

