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Genesis of Organic Computing Systems: Coupling Evolution and Learning

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Summary. Organic computing calls for efficient adaptive systems in which flexibility is not traded in against stability and robustness. Such systems have to be specialized in the sense that they are biased towards solving instances from certain problem classes, namely those problems they may face in their environment. Nervous systems are perfect examples. Their specialization stems from evolution and development. In organic computing, simulated evolutionary structure optimization can create artificial neural networks for particular environments. In this chapter, trends and recent results in combining evolutionary and neural computation are reviewed. The emphasis is put on the influence of evolution and development on the structure of neural systems. It is demonstrated how neural structures can be evolved that efficiently learn solutions for problems from a particular problem class. Simple examples of systems that "learn to learn" as well as technical solutions for the design of turbomachinery components are presented.

7.1 Introduction

Technical systems that continuously adapt to a changing natural environment and act (quasi-) autonomously have not been designed so far. Several fundamental challenges have to be met. First, more flexibility is required on the software and possibly even on the hardware level. Second, this flexibility must not be traded in against system stability and robustness. Minimal performance must be guaranteed under all circumstances and degradation must be gradual and controlled. Third, the system must be expandable and sustainable.

Biological neural systems usually have such properties while their technical counterparts do not yet meet these requirements. Nevertheless, we believe that artificial neural networks (NNs) provide a computing paradigm whose potential has not yet been fully exploited. Our approach to address the abovementioned challenges and to tap the potential of artificial neural systems is

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to tune them for particular classes of problems and particular patterns of processing.

In nature, such specialization stems from evolution and development. Both design and shape structures ready to accommodate learning and self-organizing processes, which we see as the driving forces behind the capability of neural systems, see figure 7.1. We think that understanding the biological "design techniques" for nervous systems – evolution, development, and learning – paves the way for the design of artificial adaptive systems competitive with humans.

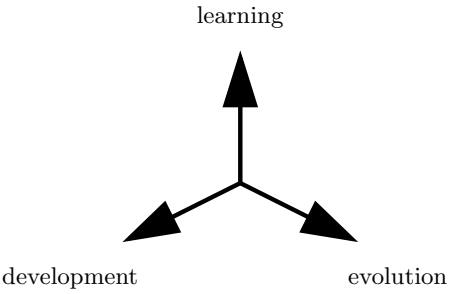

Fig. 7.1. Dimensions of natural design.

When designing adaptive systems, appropriate specialization (bias) and invariance properties are important, partially conflicting objectives. The "No-free-lunch theorems" for learning and optimization imply that it is fruitless to try to build universal adaptive systems. All systems have to be biased towards particular problem classes. This bias can be induced by evolved structures, on which learning and self-organizing processes operate. In this chapter, we review trends and recent results in combining evolutionary and neural computation. We will highlight synergies between the two fields beyond the standard examples and emphasize the influence of evolution and development on the structure of neural systems for the purpose of adaptation. We demonstrate how neural structures can be evolved that efficiently learn particular problem classes. We present simple examples of systems that "learn to learn" as well as technical solutions for the design of turbomachinery components.

The next section provides some background in artificial NNs and evolutionary algorithms (EAs). We put an emphasis on theoretical limitations and perspectives of these computing paradigms. We briefly describe simple NNs based on integrate-and-fire neurons, introduce EAs in the framework of stochastic search, and summarize the No-free-lunch theorems for learning and optimization. In section7.3, we discuss evolutionary structure optimization of neural systems and review some more recent trends in combining EAs and neural systems. Finally, we demonstrate how neural structures can be evolved that efficiently learn solutions for problems from a particular problem class.

7.2 Background

In this section, we provide short introductions to NNs and EAs and review "No-free-lunch" results for learning and optimization.

7.2.1 Neural computation

In the following, we briefly introduce basic ideas of NNs on the basis of the simple *rate-coded leaky integrator neuron* model. A detailed introduction to the broad field of neural computation is far beyond the scope of this article. A good starting point for reading is [5], recommendable introductory books on NNs for technical applications are [8, 33, 9] and on modeling nervous systems [17].

Neural systems can be described on different levels of abstraction. Many models, including those usually adopted for technical applications, can be derived from the *leaky integrator neuron*. This model is based on the assumption that the basic units of computation in nervous systems are single neurons. A model neuron i is situated in time t and its state is described by the *membrane potential* $u_i(t)$ governed by the differential equation

$$\tau_i \frac{\partial u_i(t)}{\partial t} = -u_i(t) + \sum_j w_{ij}\sigma_j[u_j(t)] + \sum_k w'_{ik}s_k(t) + \theta_i$$

with time constant τ_i. The neuron computes a weighted linear sum of the inputs it receives (see [55] for a review of more detailed models of single neurons). The first sum runs over all neurons j providing input to i, the second over all external inputs $s_k(t)$, which are gathered in the vector $\mathbf{s}(t)$, to the system. The weights w_{ij} and w'_{ik} describe the strengths of the synaptic connections. In the absence of input the membrane potential relaxes to the resting level θ_i. It is assumed that the only communication in a network of these units is through spikes of electrical activity traveling between the neurons. A neuron emits a spike when its membrane potential exceeds a certain threshold. Real spikes are discrete events, but in the model a rate code describing the average spiking frequency is assumed to capture the essence of the signals. This rate can either be viewed as an ensemble average across a population of neurons with the same properties, or as the frequency of spikes of a single neuron in some time interval. The activation function σ_i, which is usually sigmoidal (i.e., nondecreasing and bounded), maps the membrane potential u_i to the corresponding spiking frequency. Forward Euler approximation, $\partial u_i(t)/\partial t \approx (u_i(t + \Delta t) - u_i(t))/\Delta t$, with $\Delta t = \tau_i = 1$, leads to the basic discrete-time equation $u_i(t + 1) = \sum_j w_{ij}\sigma_j[u_j(t)] + \sum_k w'_{ik}s_k(t) + \theta_i$.

The structure or architecture of the NN can be described by a graph in which the nodes correspond to the neurons and there is an edge from i to j if neuron j gets input from neuron i. If the network graph contains no

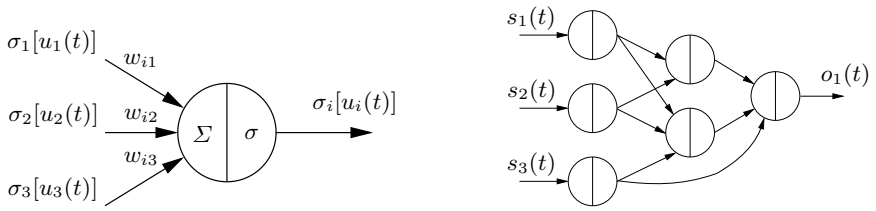

Fig. 7.2. Simple computational model of a single neuron, left, and a neural network graph, right.

cycles, we speak of a feed-forward NN. If we number the neurons such that node i only receives input from units j with $j < i$ and update the neurons in increasing order, the discrete-time network equation can be written as a static function $u_i(\mathbf{s}) = \sum_{j<i} w_{ij}\sigma_j[u_j] + \sum_k w'_{ik}s_k + \theta_i$. Often, some of the neurons are dedicated *output neurons* whose spike rates are gathered in the vector $\mathbf{o}(t)$ and the neural system can be viewed as a functional mapping input sequences $\mathbf{s}(t)$ to output sequences $\mathbf{o}(t)$. In case of a feed-forward NN, the mapping reduces to a static function assigning an output $\mathbf{o}$ to an input $\mathbf{s}$, see figure 7.2.

Models of NNs based on leaky integrator neurons, in principle exhibit universal approximation and computation properties under mild assumptions (see, e.g., [87, 89, 91]). However, the general question of how to design an appropriate neural system efficiently for a given task remains open and complexity theory reveals the need for using heuristics (see, e.g., [88]) — here these heuristics are the major organization principles of biological NNs, evolution, development, and learning.

Supervised learning of an NN means adapting the weights w_{ij}, w'_{ik} such that, given some input $\mathbf{s}(t)$, the output neurons show a predefined behavior $\mathbf{y}(t)$, which is described by sample (training) input-output sequences. A feed-forward NN learns a static function h based on sample input-output patterns $\{(\mathbf{s}_1, \mathbf{y}_1), \ldots, (\mathbf{s}_\ell, \mathbf{y}_\ell)\}$. This is usually done by gradient-based minimization of the (squared) differences between the targets $\mathbf{y}_i$ and the corresponding outputs $\mathbf{o}_i$ of the NN given the input $\mathbf{s}_i$. The ultimate goal is not to simply memorize the training patterns, but to find a statistical model for the underlying relationship between input and output data. Such a model will generalize well, that is, it will make good predictions for cases other than the training patterns. Therefore, a critical issue is to avoid overfitting during the learning process: The NN should just fit the signal and not the noise. This is usually achieved by restricting the effective complexity of the network, for example by regularization of the learning process [3].

In the context of feed-forward NNs, generalization can for example be formalized in the framework of statistical learning theory as follows. Let the goal be to learn a function from some input space S to some output space Y and

let $h : S \to Y$ be the function realized by the NN. Based on some input-output patterns drawn independently from the same distribution P on $S \times Y$, the goal of generalization is to minimize $\int_{S\times Y} P(\mathbf{s},\mathbf{y})L(h(\mathbf{s}),\mathbf{y})\mathrm{d}\mathbf{s}\,\mathrm{d}\mathbf{y}$, where $L : Y \times Y \to \mathbb{R}_0^+$ denotes a loss function. The distribution P is usually unknown and defines the learning problem at hand. The value $L(a, b)$ quantifies the cost or regret of predicting a instead of b and returns zero if its arguments are equal. For example when learning a one-dimensional real-valued function, $S = Y = \mathbb{R}$ and $L(a, b) = (a - b)^2$ is a typical choice.

In this chapter, we focus on the architecture of feed-forward neural networks, however, most of our findings and discussions apply equally well to recurrent neural systems, which also have been used successfully in applications in the past (in particular for time series prediction, e.g., [93, 61]).

7.2.2 Evolutionary computation

Evolutionary algorithms can be regarded as a special class of global random search algorithms. Let the search problem under consideration be described by a quality function $f : \mathcal{G} \to \mathcal{Y}$ to be optimized, where $\mathcal{G}$ denotes the search space (i.e., the space of candidate solutions) and $\mathcal{Y}$ the (at least partially) ordered space of cost values. The general global random search scheme can be described as follows:

① Choose a joint probability distribution $P_{\mathcal{G}^\lambda}^{(1)}$ on $\mathcal{G}^\lambda$. Set $t \leftarrow 1$.
② Obtain λ points $\mathbf{g}_1^{(t)}, \ldots, \mathbf{g}_\lambda^{(t)}$ by sampling from the distribution $P_{\mathcal{G}^\lambda}^{(t)}$. Evaluate these points using f.
③ According to a fixed (algorithm dependent) rule construct a new probability distribution $P_{\mathcal{G}^\lambda}^{(t+1)}$ on $\mathcal{G}^\lambda$.
④ Check whether some stopping condition is reached; if the algorithm has not terminated, substitute $t \leftarrow t + 1$ and return to step ②.

Random search algorithms can differ fundamentally in the way they describe (parameterize) and alter the joint distribution $P_{\mathcal{G}^\lambda}^{(t)}$, which is typically represented by a semi-parametric model. The scheme of a canonical EA is shown in figure 7.3. In evolutionary computation, the iterations of the algorithm are called *generations*. The search distribution of an EA is given by the *parent population*, the *variation operators*, and the *strategy parameters*. The parent population is a multiset of μ points $\tilde{\mathbf{g}}_1^{(t)}, \ldots, \tilde{\mathbf{g}}_\mu^{(t)} \in \mathcal{G}$. Each point corresponds to the *genotype* of an *individual*. In each generation, λ *offspring* $\mathbf{g}_1^{(t)}, \ldots, \mathbf{g}_\lambda^{(t)} \in \mathcal{G}$ are created by the following procedure: Individuals for reproduction are chosen from $\tilde{\mathbf{g}}_1^{(t)}, \ldots, \tilde{\mathbf{g}}_\mu^{(t)}$. This is called *mating selection* and can be deterministic or stochastic (where the sampling can be with or without replacement). The offspring's genotypes result from applying variation operators to these selected parents. Variation operators are deterministic or partially stochastic mappings from $\mathcal{G}^k$ to $\mathcal{G}^l$, $1 \le k \le \mu, 1 \le l \le \lambda$. An operator with $k = l = 1$

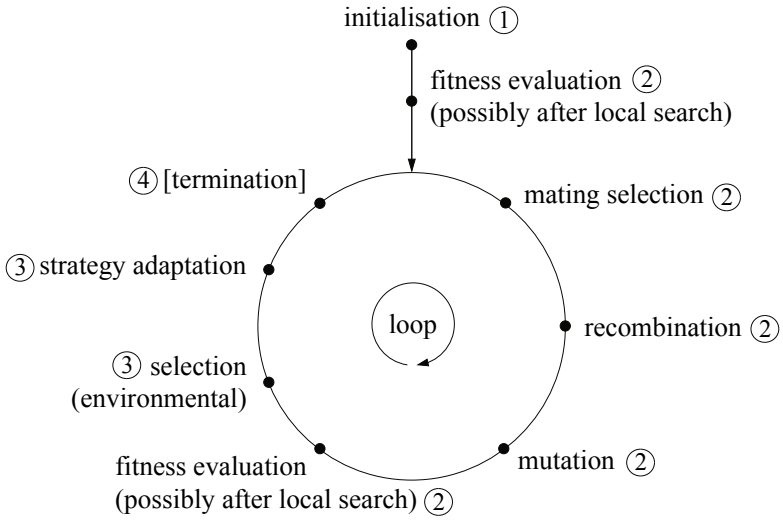

Fig. 7.3. Basic EA loop. The numbers indicate the corresponding steps in the random search scheme. When optimizing adaptive systems, the local search usually corresponds to some learning process.

is called *mutation*, whereas *recombination* operators involve more than one parent and can lead to more than one offspring. Multiple operators can be applied consecutively to generate offspring. For example, an offspring $\mathbf{g}_i^{(t)}$ can be the product of applying recombination $o_{\text{rec}} : \mathcal{G}^2 \to \mathcal{G}$ to two randomly selected parents $\tilde{\mathbf{g}}_{i_1}^{(t)}$ and $\tilde{\mathbf{g}}_{i_2}^{(t)}$ followed by mutation $o_{\text{mut}} : \mathcal{G} \to \mathcal{G}$, that is, $\mathbf{g}_i^{(t)} = o_{\text{mut}}\left(o_{\text{rec}}\left(\tilde{\mathbf{g}}_{i_1}^{(t)}, \tilde{\mathbf{g}}_{i_2}^{(t)}\right)\right)$. Evolutionary algorithms allow for incorporation of *a priori* knowledge about the problem by using tailored variation operators combined with an appropriate encoding of the candidate solutions.

Let $P_{\mathcal{G}^\lambda}^{(t)}(\mathbf{g}_1^{(t)}, \ldots, \mathbf{g}_\lambda^{(t)}) = P_{\mathcal{G}^\lambda}\left(\mathbf{g}_1^{(t)}, \ldots, \mathbf{g}_\lambda^{(t)} \,|\, \tilde{\mathbf{g}}_1^{(t)}, \ldots, \tilde{\mathbf{g}}_\mu^{(t)}; \boldsymbol{\theta}^{(t)}\right)$ be the probability that parents $\tilde{\mathbf{g}}_1^{(t)}, \ldots, \tilde{\mathbf{g}}_\mu^{(t)}$ create offspring $\mathbf{g}_1^{(t)}, \ldots, \mathbf{g}_\lambda^{(t)}$. This distribution is additionally parameterized by some *external strategy parameters* $\boldsymbol{\theta}^{(t)}$, which may vary over time. In some EAs, the offspring are created independently of each other based on the same distribution. In this case, the joint distribution $P_{\mathcal{G}^\lambda}^{(t)}$ can be factorized as $P_{\mathcal{G}^\lambda}^{(t)}(\mathbf{g}_1^{(t)}, \ldots, \mathbf{g}_\lambda^{(t)}) = P_{\mathcal{G}}^{(t)}(\mathbf{g}_1^{(t)}) \cdot \ldots \cdot P_{\mathcal{G}}^{(t)}(\mathbf{g}_\lambda^{(t)})$.

Evaluation of an individual corresponds to determining its fitness by assigning the corresponding cost value given by the quality function f. Evolutionary algorithms can — in principle — handle optimization problems that are non-differentiable, non-continuous, multimodal, and noisy. They are easy to parallelize by distributing the fitness evaluations of the offspring. In single-

objective optimization, we usually have $\mathcal{Y} \subset \mathbb{R}$, whereas in multi-objective optimization, see section 7.3.2.1, vector-valued functions (e.g., $\mathcal{Y} \subset \mathbb{R}^k, k > 1$) are considered. In co-evolution (see section 7.3.2.2), individuals interact to affect each other's adaptations. Therefore, the fitness values are not determined for each individual in isolation, but in the context of the current population (i.e., a more appropriate description of fitness assignment is $f : \mathcal{G}^\lambda \to \mathcal{Y}^\lambda$ or even $f : \mathcal{G}^{\lambda+\mu} \to \mathcal{Y}^{\lambda+\mu}$ if the parents are also involved in the fitness calculation). The interaction of individuals may be competitive or cooperative. As the fitness function is not fixed, co-evolution allows for "bootstrapping" the evolutionary process and "open-ended" evolution.

Updating the search distribution corresponds to *environmental selection* and sometimes additional *strategy adaptation* of external strategy parameters $\boldsymbol{\theta}^{(t+1)}$. The latter is extensively discussed in the context of optimization of NNs in [40, 45]. A selection method chooses μ new parents $\tilde{\mathbf{g}}_1^{(t+1)}, \ldots, \tilde{\mathbf{g}}_\mu^{(t+1)}$ from $\tilde{\mathbf{g}}_1^{(t)}, \ldots, \tilde{\mathbf{g}}_\mu^{(t)}$ and $\mathbf{g}_1^{(t)}, \ldots, \mathbf{g}_\lambda^{(t)}$. This second selection process is called environmental selection and may be deterministic or stochastic. Either the mating or the environmental selection must be based on the objective function values of the individuals and must prefer those with better fitness — this is the driving force of the evolutionary adaptation process.

It is often argued that evolutionary optimization is not well understood theoretically — ignoring the tremendous progress in EA theory during the last years. Although there are only a few results for general settings (e.g., convergence [76]), there exist rigorous expected runtime analyses of simplified algorithms on restricted, but important classes of optimization problems, see [46, 20] and references therein. The article [7] provides a good starting point for reading about EA theory.

7.2.3 The need for specialization: No-free-lunch

It is not only intuitive, but also proven that it is not possible to design an universal adaptive system that outperforms other systems across all possible problems. This is formally expressed by the No-free-lunch (NFL) theorems going back to the work of Wolpert and Macready [104, 105]. Coarsely speaking, the NFL theorems for learning state that without an assumption of how the past (training data) is related to the future (test data), prediction is impossible. In other words, without an a priori restriction of the possible phenomena that are expected, it is impossible to generalize and thus no algorithm is superior to another. Even worse, any consistent algorithm (i.e., any algorithm converging to the Bayes optimal classifier almost surely when the number of training patterns, drawn independently from the distribution describing the problem, approaches infinity) can have arbitrarily poor behavior when given a finite, incomplete training set [104, 19, 10].

These results carry over to general search and optimization algorithms. The NFL theorem for optimization formalizes that averaged over the set $\mathcal{F}$

of all possible objective functions defined between a finite search space $\mathcal{X}$ and a finite set $\mathcal{Y}$ of cost values all optimization algorithms have the same performance. It is assumed that the algorithms never visit a search point twice and that the performance measure just depends on the objective function values of the visited search points [20, 105, 82, 42, 43, 106]. More generally, the following holds for any probability distribution P over $\mathcal{F}$. If and only if $\mathcal{F} = \bigcup_i \mathcal{F}_i$, every $\mathcal{F}_i$ is closed under permutation, and $f, g \in F_i$ implies that f and g have the same probability $P(f) = P(g)$ to be the objective function, then all optimization algorithms have the same performance averaged over $\mathcal{F}$ w.r.t. P [43]. Closure under permutation of a set $\mathcal{F}_i$ means that for every bijective function $\pi : \mathcal{X} \to \mathcal{X}$ it holds that $f \in \mathcal{F}_i$ implies $f \circ \pi \in \mathcal{F}_i$. These assumptions for an NFL result to hold are rather strict, and fortunately problem classes relevant in practice are likely to violate them [42, 43].

Nonetheless, only if we consider restricted problem classes, in which the assumptions of the NFL theorems are not fulfilled, we can design efficient adaptive systems. It is important to make this bias towards a problem class explicit in the design process. In nature, such a bias stems from the evolved structures on which learning and self-organizing processes operate. In organic computing, simulated evolutionary structure optimization can create systems biased towards relevant problem classes.

7.3 Evolutionary computation and neural systems

Both artificial evolution and artificial neural systems have long histories, which in many respects resemble each other. In their beginnings, both were met with considerable skepticism from the biological as well as from the technological communities. For the first, their simplifications and abstractions meant throwing over board years of carefully accumulated details on how biological systems operate, develop, learn, and evolve. For the second the new type of distributed, stochastic, and nonlinear processing was equally hard to accept. During their maturation both fields met a couple of times, but not as often as one might expect bearing in mind that their philosophies to extract principles of biological information processing and apply them to technical systems are so similar.

Although not directly aimed at the formation of neural systems, the design of intelligent automata was among the earliest applications of EAs and may be traced back to the 1950s, see [22]. However, it took another 30 years until first papers were published describing explicitly the application of EAs to NNs and in this context — albeit more hesitantly — to learning [52, 63]. Then the subject quickly received considerable interest and several articles were published in the early nineties concentrating on optimization of both the network architecture and its weights. Although nowadays NNs and EAs are used frequently and successfully together in a variety of applications, the desired

breakthrough, that is, the evolution of neural systems showing *qualitatively* new behavior, has not been reached yet. The complexity barrier may have been pushed along but it has not been broken down. Nevertheless, many important questions on the architecture, (e.g., modularity) the nature of learning, (e.g., nature vs. nurture) and the development of neural systems (e.g., interactions between levels of adaptation) have been raised and important results have been obtained.

There are still only few works connecting current brain research with evolutionary computation, but first attempts have been promising, as we will see in section 7.3.2.5. Here, on a more general note, we argue that combining evolutionary development with brain science is more than just optimizing models of biological neural systems. The brain is a result of the past as much as of the present. That means that learning (the present) can only operate on an appropriate structure (the past). The current structure reflects its history as much as its functionality. Flexibility and adaptability of the brain are based on its structural organization, which is the result of its ontogenetic development. The brain is not one design but many designs; it is like a cathedral where many different parts have been added and removed over the centuries. However, not all designs are capable of such continuous changes and the fact that the brain is, is deeply rooted in its structural organization.

In this section, we discussed selected aspects of combining neural and evolutionary computing. More comprehensive surveys, all having slightly different focuses, can be found in [65, 71, 83, 107].

7.3.1 Structure optimization of adaptive systems

Although NNs are successfully applied to support evolutionary computation (see section 7.4.2), the most prominent combination of EAs and NNs is evolutionary optimization of adaptive neural systems.

In general, the major components of an adaptive system can be described by a triple $(\mathcal{S}, \mathcal{A}, \mathcal{D})$, where $\mathcal{S}$ stands for the structure or architecture of the adaptive system, $\mathcal{A}$ is a learning algorithm that operates on $\mathcal{S}$ and adapts flexible parameters of the system, and $\mathcal{D}$ denotes sample data driving the adaptation. We define the *structure* as those parts of the system that cannot be changed by the learning/self-adaptation algorithm. Given an adaptation rule $\mathcal{A}$, the structure $\mathcal{S}$ determines

- the set of solutions that can be realized,
- how solution changes given new stimuli/signals/data, partial failure, noise, etc.,
- the neighborhood of solutions (i.e., distances in solution space),
- bias (specialization) and invariance properties.

Learning of an adaptive system can be defined as goal-directed, data-driven change of its behavior. Examples of learning algorithms for technical

NNs include gradient-based heuristics (see section 7.2.1) or quadratic programming. Such "classical" optimization methods are usually considerably faster than pure evolutionary optimization of these parameters, although they might be more prone to getting stuck in local minima. However, there are cases where "classical" optimization methods are not applicable, for example when the neural model or the objective function is non-differentiable (e.g., see section 7.3.2.2). Then EAs for real-valued optimization provide a means for adjusting the NN parameters. Still, the main application of evolutionary optimization in the field of neurocomputing is adapting the structures of neural systems, that is, optimizing those parts that are not altered by the learning algorithm. Both in biological and technical neural systems the structure is crucial for the learning behavior — the evolved structures of brains are an important reason for their incredible learning performance: "development of intelligence requires a balance between innate structure and the ability to learn" [6]. Hence, it appears consequential to apply evolutionary methods to structure adaptation of neural systems for technical applications, a task for which usually no efficient "classical" methods exist.

A prototypical example of evolutionary optimization of a neural architecture on which a learning algorithm operates is the search for an appropriate topology of a multi-layer perceptron NN, see [103, 36, 29] for some real-world applications. Here, the search space ultimately consists of graphs, see section 7.2.1. When using EAs to design NN graphs, the key questions are how to encode the topologies and how to define variation operators that *act* on this representation. In the terminology of section 7.2.2, operators and representation both determine the search distribution and thereby the neighborhood of NNs in the search space. Often an intermediate space, the phenotype space $\mathcal{P}$, is introduced in order to facilitate the analysis of the problem and of the optimization process itself. The fitness function can then be written as $f = f' \circ \phi$, where $\phi : \mathcal{G} \to \mathcal{P}$ and $f' : \mathcal{P} \to \mathcal{Y}$. The definition of the phenotype space is to a certain degree arbitrary. The same freedom exists in evolutionary biology [60] and is not restricted to EAs. The probability of a certain phenotype $p \in \mathcal{P}$ to be created from a population of phenotypes strongly depends on the representation and the variation operators. When the genotype-phenotype mapping ϕ is not injective, we speak of neutrality, which may considerably influence the evolutionary process (see [41] for an example in the context of NNs). We assume that $\mathcal{P}$ is equipped with an extrinsic (i.e., independent of the evolutionary process) metric or at least a consistent neighborhood measure, which *may* be defined in relation to the function of the individual. In the case of NNs, the phenotype space is often simply the space of all possible connection matrices of the networks. Representations for evolutionary structure optimization of NNs have often been classified in "direct" and "indirect" encodings. Roughly, a direct encoding or representation is one where (intrinsic) neighborhood relations in the genotype space (induced by $P_{\mathcal{G}^\lambda}$) broadly correspond to extrinsic distances of the corresponding phenotypes. Note that such a classification only makes sense once a phenotype

space with an extrinsic distance measure has been defined and that it is only valid for this particular definition. (This point has frequently been overlooked because of the implicit agreement on the definition of the phenotype space, e.g., the graph space equipped with a graph editing distance). This does not imply that both spaces are identical. In an indirect encoding the genotype usually encodes a rule, a program or a mapping to build, grow or develop the phenotype. Such encodings foster the design of large, modular systems. Examples can be found in [54, 32, 25, 83, 84].

7.3.2 Trends in combining EAs and neural computation

In the following, we review some more recent trends in combining neural and evolutionary computing. Needless to say that such a collection is a subjective, biased selection.

7.3.2.1 Multi-objective optimization of neural networks

Designing a neural system usually requires optimization of several, often conflicting objectives. This includes coping with the bias-variance dilemma or trading classification speed against accuracy in real-time applications. Although the design of neural systems is obviously a multi-objective problem, it is usually tackled by aggregating the objectives into one scalar function and applying standard methods to the resulting single-objective task. However, this approach will in general fail to find all desired solutions [16]. Furthermore, the aggregation weights have to be chosen correctly in order to obtain the desired result. In practice, it is more convenient to make the trade-offs between the objectives explicit (e.g., to visualize them) after the design process and select from a diverse set of systems the one that seems most appropriate. This can be realized by "true" multi-objective optimization (MOO, [48]). The MOO algorithms approximate the set of Pareto-optimal tradeoffs, that is, those solutions that cannot be improved in any objective without getting worse in at least one other objective. From the resulting set of systems the final solution can be selected after optimization. There have been considerable advances in MOO recently, which can now be incorporated into machine learning techniques. In particular, it was realized that EAs are very well suited for multi-criterion optimization and they have become the MOO methods of choice in the last years [13, 18]. Recent applications of evolutionary MOO to neural systems address the design of multi-layer perceptron NNs [1, 2, 49, 103, 29, 11, 48] and support vector machines (SVMs) [39, 97].

7.3.2.2 Reinforcement learning

In the standard reinforcement learning (RL) scenario [100, 95, 74], an agent perceives stimuli from the environment and decides which action to take based

on its policy. Influenced by the actions, the environment changes its state and possibly emits reward signals. The reward feedback may be sparse, unspecific, and delayed. The goal of the agent is to adapt its policy, which may be represented by (or be based on) a NN, such that the expected reward is maximized. The gradient of the performance measure with respect to NN parameters can usually not be computed (but estimated in case of stochastic policies, e.g., see [96, 56]).

Evolutionary algorithms have proved to be powerful and competitive methods for solving RL problems [64, 38, 72]. The recent success of evolved NNs in game playing [12, 23, 59, 94] demonstrates the potential of combining NNs and evolutionary computation for RL. The possible advantages of EAs compared to standard RL methods are that they allow — in contrast to the common temporal difference learning methods — for direct search in the space of (stochastic as well as deterministic) policies. Furthermore, they are often easier to apply and more robust with respect to the tuning of the meta-parameters (learning rates, etc.). They can be applied to non-differentiable function approximators and even optimize their underlying structure.

Closely related is the research area of evolutionary robotics devoted to the evolution of "embodied" neural control systems [66, 57, 70, 101]. Here promising applications of the principle of co-evolution can be found.

7.3.2.3 Evolving network ensembles

Ensembles of NNs that cooperatively solve a given task can be preferable to monolithic systems. For example, they may allow for task decomposition that is necessary for efficiently solving a complex problem and they are often easier to interpret [85]. The population concept in EAs appears to be ideal for designing neural network ensembles, as, for example, demonstrated for classification tasks in [58, 11]. In the framework of decision making and games, Mark et al. [62] developed a combination of NN ensembles and evolutionary computation. Two ensembles are used to predict the opponent's strategy and to optimize the own action. Using an ensemble instead of a single network ensures to be able to maintain different opponent experts and counter-strategies in parallel. The EA is used to determine the optimal input for the two network ensembles. Ensembles of networks have also turned out a superior alternative to single NNs for fitness approximation in evolutionary optimization. In [51] network ensembles have been optimized with evolution strategies and then used as metamodels in an evolutionary computation framework. Beside the increase in approximation quality an ensemble of networks has the advantage that the fidelity of the networks can be estimated based on the variance of the ensemble.

7.3.2.4 Optimizing kernel methods

Adopting the extended definition of structure as that part of the adaptive system that cannot be optimized by the learning algorithm itself, model selection

of kernel-based methods is a structure optimization problem. For example, choosing the right kernel for an SVM [14, 15, 81] is important for its performance. When a parameterized family of kernel functions is considered, kernel adaptation reduces to finding an appropriate parameter vector. These "hyperparameters" are usually determined by grid search, which is only suitable for the adjustment of very few parameters, or by gradient-based approaches. When applicable, the latter methods are highly efficient albeit susceptible to local optima. Still, the gradient of the performance criterion w.r.t. the hyperparameters can often neither be computed nor accurately approximated. This leads to growing interest in applying EAs to model selection of SVMs. In [26, 77, 39, 97], evolution strategies (i.e., EAs tailored for real-valued optimization) were proposed for adapting SVM hyperparameters, in [21, 27] genetic algorithms (EAs that represent candidate solutions as fixed-length strings over a finite alphabet) were used for SVM feature selection.

7.3.2.5 Computational neuroscience and brain-inspired architectures

There are only a few applications of evolutionary computation in brain science [4, 80, 92, 44, 75], although evolutionary "analysis by synthesis" guided by neurobiological knowledge may be a powerful tool in computational neuroscience. The challenge is to force artificial evolution to favor solutions that are reasonable from the biological point of view by incorporating as much neurobiological knowledge as possible in the design process (e.g., by a deliberate choice of the basic system structure and constraints that ensure biological plausibility).

In the field of brain-inspired vision systems [28, 102] EAs have been used to optimize the structure of the system (i.e., feature banks or hierarchical layers) and to determine a wide variety of parameters. Evolutionary algorithms have been successfully applied to the Neocognitron structure [98, 68, 86], which was one of the first hierarchical vision systems based on the structure of its biological counterpart [28]. More recent work employed evolution strategies to optimize the nonlinearities and the structure of a biologically inspired vision network, which is capable of performing a complex 3D real world object classification task [78, 79]. The authors used evolutionary optimization with direct encoding that performed well in an 1800-dimensional search space. In a second experiment evolutionary optimization was successfully combined with local unsupervised learning based on a sparse representation. The resulting architecture outperformed alternative approaches.

7.4 Networks that learn to learn

The ability to learn (online) is one of the most distinguishing features of artificial NNs. The idea behind the "learn to learn" concept discussed in this section

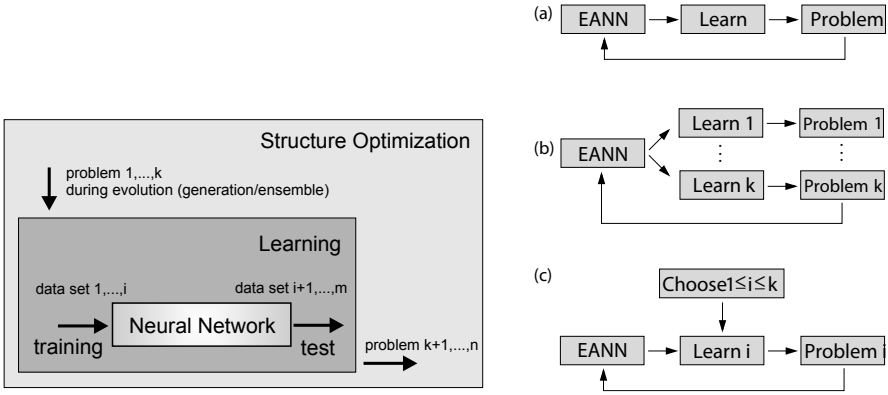

Fig. 7.4. Evolutionary structure optimization for problem classes. Left: Methods to achieve first and second order generalization in neural network learning and evolution (structure optimization). Right: Standard neural network learning (a), parallel switching between problems (b) and sequential switching (c). EANN denotes a neural network optimized by an evolutionary algorithm.

is that the goal of evolutionary NN structure optimization should be the ability to efficiently learn new related problems during operation, see [34, 37]. Here "efficient" means fast and based on incomplete data. The term "new related problems" is more difficult to define. The problems must have some common structure that can be captured by the EA and reflected in the NN architecture. Learning a different problem class goes beyond standard generalization. The latter means generalizing from a finite set of training samples to arbitrary samples drawn from the same distribution as, for example, formally defined at the end of section 7.2.1. Facing a different problem from the same class means that the underlying distribution has changed while belonging to the set of distributions which define the class and which have some common features that can be represented by the structure.

Therefore we speak of "second order generalization" for the ability to efficiently switch between problems, see figure 7.4 (left). In the notation introduced by Thrun and Pratt [99], this ability belongs to the area of representations and functional decompositions. However, in the evolutionary approach, this functional decomposition is self-organized during the evolutionary process. There are basically two different ways in which second order generalization can be achieved and used: the parallel and the sequential way. In figure 7.4 (right, a) the standard approach to learn one problem with an NN is shown. In part (b), the parallel approach is shown. The network is optimized during evolution in order to learn one of a number of possible problems. The actual decision is made after the network's structure has been fixed by evolutionary search. However, during the search the network's structure must be optimized in order to cope with any of the possible problems. Thus, each structure is ap-

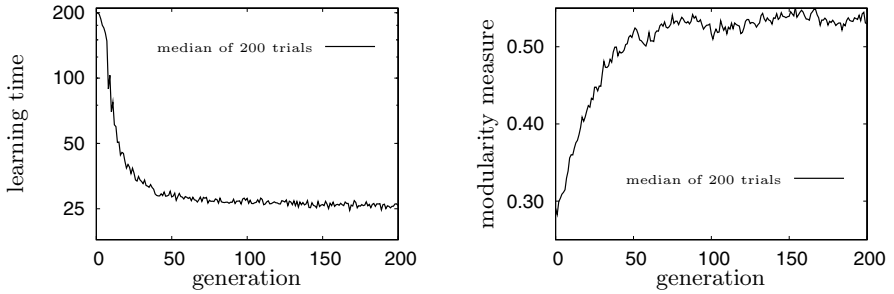

Fig. 7.5. Evolving networks that "learn to learn." During evolution the network structures adapt to a special problem class. This specialization leads to a reduced time for learning a new instance of the problem class, see left plot. Since the class in this example consists of separable problems, the degree of modularity of the network structures increases over time, as shown in the right plot (cf. [35]).

plied to all problems (or a random subset of problems of the respective class). For each problem the weights are newly initialized. The fitness of the network is determined by the mean (or median or weighted sum) of the networks' individual performances. In figure 7.4 (right, c), the network has to learn a number of problems one after the other during operation. The network's structure has been optimized in such a way that switching from problem to problem can be achieved most efficiently in the above sense. The weights are not randomly initialized (like in (b)), but averaged Lamarckian inheritance [36] is used to exploit information on previous problems for the next problem belonging to the same class. Again, the fitness of the network is determined by the mean (or median or weighted sum) of the networks' individual performances.

From the NFL theorems (see section 7.2.3) we conclude that adaptive systems have to be specialized towards a particular problem class to show above average performance. Second order generalization can be viewed as such a specialization.

7.4.1 Modularity

A simple example of how to build NNs that "learn to learn" was given in the study [35], where Hüsken et al. considered feed-forward NNs that had to learn binary mappings $\{0,1\}^6 \rightarrow \{0,1\}^2$ assigning target values $\mathbf{y} = (y_1, y_2)' \in \{0,1\}^2$ to inputs $\mathbf{s} = (s_1, \ldots, s_6)' \in \{0,1\}^6$. The class of mappings was restricted to those which are separable in the strict sense that y_1 only depends on the inputs $s_1, \ldots, s_3$ and y_2 only on $s_4, \ldots, s_6$. The mappings changed over time and a simple EA was employed to create feed-forward network structures that quickly learn a new instance of the problem class. The fitness of an NN structure was determined by the time needed to learn a ran-

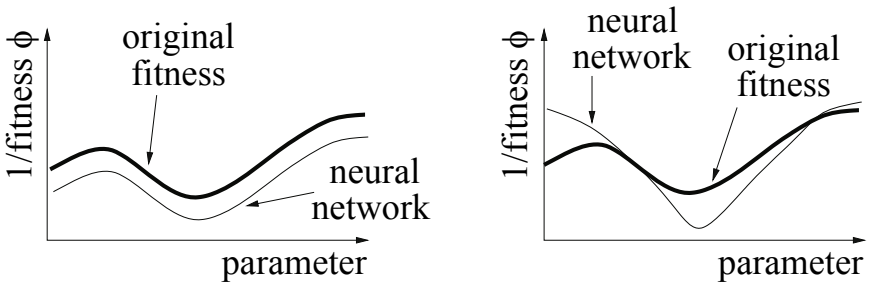

Fig. 7.6. Although the approximation errors of the neural network models are quite high, the optimization based on the approximation models leads to the desired optimum of the fitness under rank-based selection.

domly chosen problem instance, that is, the sequential switching approach depicted in figure 7.4 was used.

After a few generations, the networks adapted to the special, restricted problem class and the learning time decreased drastically, see figure 7.5. In this toy example, it is obvious that NN structures that are modular in the sense that they process the inputs $s_1, \ldots, s_3$ and $s_4, \ldots, s_6$ separately without interference are advantageous. When measuring this special kind of modularity during the course of evolution, it turned out that the modularity indeed increased, see figure 7.5, right plot.

In [53] modularity is analyzed in the context of problem decomposition and a novel modular network architecture is presented. Modularity is related to multi-network systems or ensembles for which a taxonomy is presented. A co-evolutionary framework is used to design modular NNs. The model consists of two populations, one consisting of a pool of modules and the other synthesizing complete systems by drawing elements from the first. In this framework, modules represent parts of the solution which co-operate with each other to form a complete solution. Using two artificial tasks the authors demonstrate that modular neural systems can be co-evolved. At the same time, the usefulness of modularity depends on the learning algorithm and the quality function.

7.4.2 Real-world application

Evolutionary algorithms combined with computational fluid dynamics (CFD) have been applied successfully to a large variety of design optimization problems in engineering (e.g., [90, 24, 67]). The fluid-dynamics simulations necessary to determine the quality of each design are usually computationally expensive, for example the calculation of the three-dimensional flow field around a car takes between 10-30 hours depending on the required accuracy. Therefore, metamodels or surrogates are used during the search to approximate the

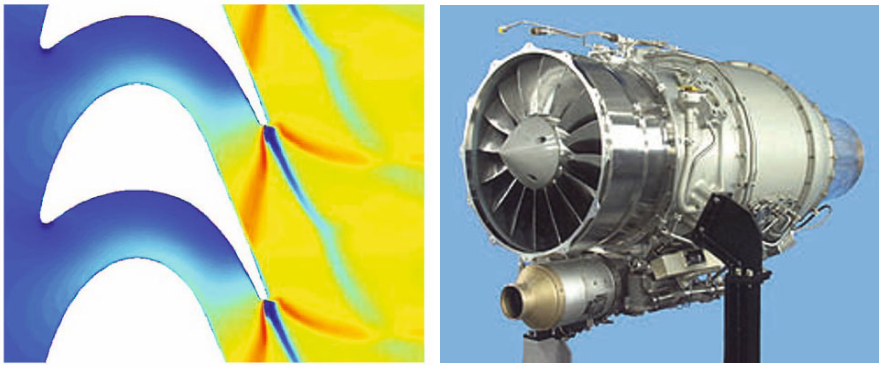

Fig. 7.7. The flow field of a turbine blade cascade for a gas turbine engine shown on the right.

results of the CFD simulations. Although first approaches to combine fitness approximation with EAs are relatively old [31], it is only in the last couple of years that the field has received wider attention, see [47] for a review. It has been revealed that the strategy to keep the update of the metamodel and the optimization process separate is not advisable, since the optimization is easily misled if the modeling quality is limited (which is often the case in practical applications). Jin et al. [50] have suggested to use the metamodel alongside the true objective function to guarantee correct convergence. Furthermore, the use of NNs as models is particularly advantageous because of their online learning ability. Thus, the approximation quality of NNs can be continuously improved during the optimization process when new CFD data is available (e.g, [50, 73, 69, 30]). It is interesting to note that the standard mean squared error measure of NNs is not necessarily the best means to determine the quality of NNs that are employed as surrogates. Figure 7.6 shows why this is the case. During evolutionary search, the absolute error of the NN is of no concern, as long as the model is able to distinguish between "good" and "bad" individuals.

7.4.2.1 Evolution of the metamodel

Neural networks that are used as metamodels during evolutionary search should have the best possible architecture for the approximation task. Therefore, EAs are employed to determine the structure of the networks offline, for example using data from previous optimization tasks. Weight adaptation is conducted during the evolutionary design optimization whenever new data are available.

This framework has been employed in [36] for the optimization of turbine blades of a gas turbine engine. The flow field around a turbine blade and the engine are shown in figure 7.7. Navier-Stokes equations with the (k-ϵ) tur-

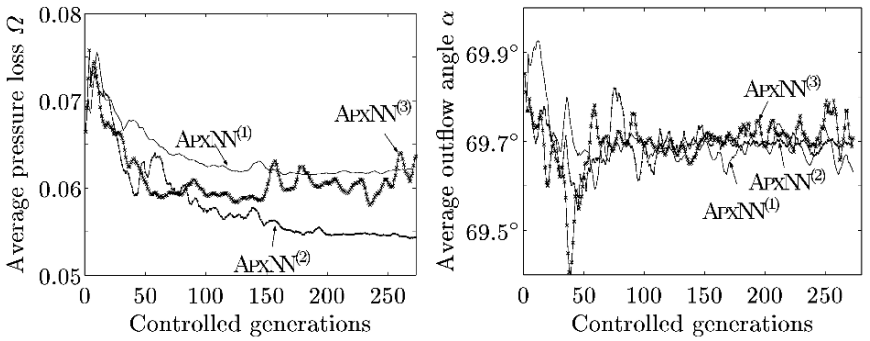

Fig. 7.8. Results normalized to the number of generations where the CFD simulations have been used. ApxNN(1) denotes a fully connected neural network, for ApxNN(2) the network structure has been evolutionarily optimized and for ApxNN(3) the network has been optimized to switch between different design domains (problem classes) most efficiently.

bulence model were used for the two dimensional CFD simulations. During optimization the pressure loss was minimized subject to a number of geometrical and functional constraints, in particular the target outflow angle α was set to 69.70 deg. The turbine blades were represented by 26 control points of non-uniform rational B-splines. The (x, y)-coordinates of the control points were optimized using a (2,11)-evolution strategy, further details can be found in [36].

The results of the optimization are given in figure 7.8. The average pressure loss and outflow angle are shown that have been reached in the evolutionary design optimization of the turbine blade. The three curves represent three different strategies to define the architecture of the NN that has been used as a metamodel during search. The model of the first type (ApxNN(1)) uses a fully connected architecture. The weights are initialized by means of offline learning, using training data collected in a comparable blade optimization trial (e.g., different initialization but the same number of control points of the spline and the same fitness function). The second type of network model (ApxNN(2)) was optimized offline with an EA using data generated in a previous optimization run. The third approach will be discussed in the next section. It is evident that the evolutionarily optimized NN structure clearly outperforms the fully connected model in the practical application.

7.4.2.2 *Learn* surrogates to *learn* CFD

We already discussed the idea to evolve the architecture of NNs not just for one specific problem but instead to optimize the network so that it is able to quickly adapt to problems belonging to one class. We can transfer this idea

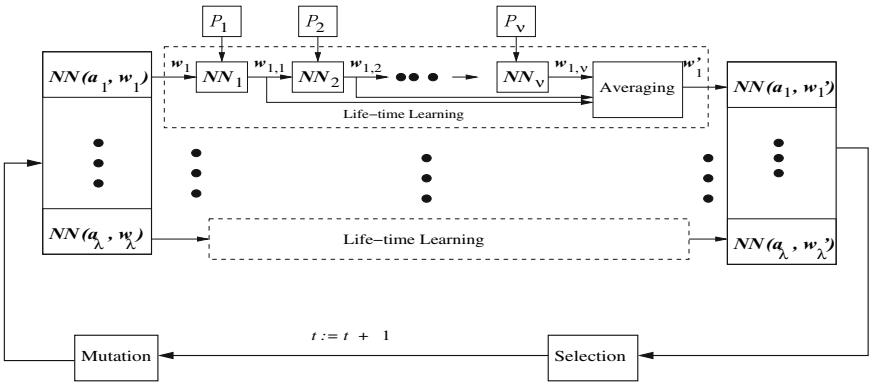

Fig. 7.9. Evolutionary structure optimization for problem classes with averaged Lamarckian inheritance, where the $\mathcal{P}_i$ denote the different problems belonging to one class and NN_i the network after learning $\mathcal{P}_i$. The symbol $w_{i,j}$ denotes the set of weights of the ith network after learning the jth problem, w_i' refers to the weights of the ith network after learning all ν problems. Averaged Lamarckian evolution is used to take the different problem characteristics into account for determining the set w_i', details can be found in [36]. The symbol a_i denotes the architecture or structure of the neural network, which is not changed during the sequential learning of problems $1 \ldots \nu$.

to the problem domain of surrogates for approximation during evolutionary design optimization by sub-dividing the CFD samples into groups (problems) belonging to one and the same class namely the approximation of CFD data for evolutionary search. This is a reasonable approach because we do not expect to evolve a network that approximates the CFD results well for the whole optimization. Instead, since the surrogate and the original CFD simulation are mixed during search, new data samples are available and the network can be adapted online. Thus, the best network is the one that is particularly well suited to continuously and quickly learn new CFD approximations during the evolutionary design optimization. In figure 7.9 the framework for the evolutionary optimization of the NN for problem classes is shown. To avoid confusion, we point out that the evolutionary optimization of the architecture is still decoupled from the evolutionary design optimization, where the best network is used as a surrogate.

The results for the network that has been evolved to quickly adapt to problems from one and the same class are shown as ApxNN(3) in figure 7.8. We observe that during the first generations ApxNN(3) scores much better than ApxNN(1) (the fully connected NN) and similar to ApxNN(2) (the network whose structure was optimized using a standard evolutionary approach to minimize the approximation error for all data offline). However, in later generations, the performance of ApxNN(3) deteriorates and becomes unstable. Although this

behavior is not yet fully understood, we believe that one reason might be the different update frequencies between the offline problem class training and the online design optimization. The update frequency denotes how often the original CFD simulation is called within a certain number of generations. As this frequency is adapted depending on the fidelity of the approximation model, it changes differently during offline structure optimization of the NN and online application of the network as a surrogate for the design optimization. Therefore, the definition of the problem class might change, which is difficult to cope with for the network.

7.5 Conclusion

Organic computing calls for adaptive systems. In order to be efficient and robust, these systems have to be specialized to certain problem classes comprising those scenarios they may face during operation. Nervous systems are perfect examples of such specialized learners and thus are prime candidates for the substrate of organic computing.

Computational models of nervous systems like artificial neural networks (NNs) have to be revisited in the light of new adaptation schemes that focus on the structure of the system and address issues like modularity, second-order generalization and learning efficiency.

At the same time, we promote the combination of evolutionary algorithms (EAs) and NNs not just because of an appealing metaphor, but also and foremost because EAs have proved to be well suited to solve many of the difficult optimization problems occurring when designing NNs, especially when higher order optimization methods cannot be applied. The field of evolutionary neural systems is expanding in many different directions as we have shown in this chapter. We have demonstrated how NNs can be evolved that are specialized to certain problem classes. Although still in its beginnings, this second order learning is not restricted to toy problems but has already found real-world technical applications.

Still, much is left to do to establish the design triangle learning–development–evolution of neural systems in such a way that they can demonstrate their full potential. Results from brain science highlight the importance of architecture and of the way the architecture is constructed during ontogenesis. Although the incorporation of evolution and development into computational neuroscience is still in its beginning, we believe that this will be a promising approach to organic computing.

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