1. Introduction

1.1 Hearing and learning

Learning is the essence of life. Throughout evolution, all species learn to adapt to a particular environment. As a consequence, organs are formed and reformed, and cellular function extends to respond to a broadening spectrum of types of energy and chemical compounds and focuses on particular stimulus configurations to respond to them in ever subtler ways. In ontogeny, chemical and spatial orientation is essential for the cells of an embryo so that it can grow all parts at the right time and at the right location. Postnatally, the radius from which an organism needs to learn expands from itself to the world surrounding it. For mammals, there appear to be two distinctly different phases of learning from the environment: early in life, or approximately before sexual maturity, and later in life, or post puberty. It has been a remarkable neuroscientific discovery that the molecular machinery employed by organism during their development is subtly but effectively modified to provide neuroplasticity later in life.

We learn to hear in ontogeny beginning at prenatal times (Peña et al., 2003), and patients receiving a cochlear implant (CI) need to learn or to re-learn hearing, depending on whether they never heard or they originally had a normal hearing experience. Both challenges require specific forms of learning. Learning implies appropriate adjustments of nervous networks to reflect differences and changes in patterns of sensory activity. Neuroscience looks for molecular, structural, and functional consequences of learning in the nervous system and refers to these changes as indicators of neuronal plasticity. To study sensory-evoked learning and plasticity we need to look at the molecular profile and cellular dynamic of neurons to identify its lasting traces in the brain.

One molecular marker associated to the initiation of sensory-evoked neuronal remodeling is the early response gene product c-Fos. Among the markers reflecting a growth-response of nerve cells and their synapses is the growth-associated protein GAP-43.
1.2 Auditory brainstem plasticity

Although earlier reports already suggested the possibility of structural plasticity in the adult mammalian brainstem (Wall & Egger, 1971; Pollin & Albe-Fessard, 1979), this mode of learning remained largely unacceptable into the 1980s. However, evidence slowly began to accumulate that structural plasticity is a general phenomenon of the adult mammalian brain, including all ages and all levels of the cerebral hierarchy. Equally important was the discovery that adult plasticity reuses many molecular mechanisms and regulatory circuits on which embryogenesis relies. However, important differences exist between juvenile and adult brain plasticity with respect to speed, breadth, and type of changes inducible by internal or external stimuli (e.g. Harris et al., 2005; 2008). At all developmental stages, neuroplasticity has many facets and includes the mutability on virtually all levels of neuronal organization, including volume growth, modifications of topographical maps, axon rerouting, dendritic outgrowth or retraction, cell death, mitosis, synaptogenesis, and molecular changes of various sorts. We finally appreciate the brain as an instrument of change.

1.3 c-Fos

1.3.1 c-Fos and learning

Immediate early genes are essential actors mediating learning-associated cellular processes and neuroplasticity. Although the immediate early gene product c-Fos has, together with phosphorylated extracellular signal-related kinase (p-ERK), been used as a marker for central sensitization, particularly in studies using nociception (Gao & Ji, 2009), its expression signifies not just electrophysiological activity but also cellular activity on different levels. Expression of c-Fos, like other early response genes, links fleeting changes of neuronal activity to lasting modifications of structure and function in the mammalian nervous system. Zuschratter et al. (1995) compared the spatial distribution of c-Fos immunoreactive neurons with the density of 2-desoxy-D-glucose (2DG) autoradiography after 1h of acoustic stimulation. While the pattern of the highest density of c-Fos labeled cells in the auditory cortex matched the peak labeling of autoradiographs, a spreading of c-Fos expression in neurons across the tonotopic maps was observed in primary auditory cortex and in the rostral and caudal fields of the auditory cortex. Corresponding to its role beyond indicating electrophysiological activity, c-Fos has been reported to be involved in the learning correlate of long term potentiation (Racaniello et al., 2010). Watanabe et al. (1996) found that homozygous mice carrying a null mutation of c-fos fail to show much of functional and structural plasticity seen in normal mice.

Auditory cortical neurons were shown to turn c-Fos positive following behaviorally significant sounds (Fichtel & Ehret, 1999; Wan et al., 2001; Geissler & Ehret, 2004). Sound recognition was compared with perception of exactly the same sound in mice. They showed that sound recognition, relying on memory, entails less but well focused Fos-positive cells in a primary auditory cortical field and significantly more labeling in higher order fields.

1.3.2 Regulation of c-Fos

The immediate early gene c-fos has, as part of its promoter, a transcription factor binding site called the cAMP response element (CRE; Ginty et al., 1994; Gispen et al., 1991). Thus, glutamate binding on NMDA receptors may activate the c-fos gene in neurons through the
calcium-dependent phosphorylation of the CRE binding protein CREB by ERK and/or CaMKIV kinase pathways. CREB phosphorylation is modulated in cochlear nucleus and superior olivary complex as a consequence of electrical stimulation in the cochlea (Illing & Michler, 2001). Apart from CRE, the c-fos promoter contains a sis inducible element (SIE), a serum response element (SRE), and an activator protein-1 (AP-1)-like sequence (FAP) (Herdegen & Leah, 1998). Besides its acting on the c-fos promoter, p-CREB may also affect expression of other genes containing a CRE site. Among them are the genes coding for the brain derived neurotrophic factor BDNF (Tao et al., 1998; Xu et al., 2000) and the early growth response protein-1 (Egr-1) (Sakamoto et al., 1991; Schwachtgen et al., 2000), another immediate-early gene product which is also known as Krox-24, Zif268, Zenk, NGFI-A, and Tis8. Apart from its CRE site, the promoter of the Egr-1 gene contains a SRE site and an AP-1 site (Schwachtgen et al., 2000; Weber & Skene, 1998). Competing with the activating transcription factor-2 (ATF-2; van Dam & Castellazzi, 2001), the protein c-Fos may dimerize with still another immediate-early gene product, c-Jun, to constitute a powerful transcription factor complex, called AP-1. The AP-1 factor, in turn, differentially triggers the expression of a large number of genes in a variety of functional aspects (Wisdom, 1999), with ATF-2:Jun and Fos:Jun complexes having differential binding preferences for heptameric or octameric AP-1 binding sites (van Dam and Castellazzi, 2001). Among the genes controlled by AP-1 are genes coding for the basic fibroblast growth factor bFGF (Shibata et al., 1991) and the growth and plasticity associated protein GAP-43 (Nedivi et al., 1992; Weber & Skene, 1998). Deficiency in ATF-2, in turn, leads to neurodegeneration of subsets of somatic and visceral motorneurons of the brainstem (Ackermann et al., 2011). There are conditions under which c-Fos may appear in glial cells (Edling et al., 2007).

Looking further upstream of c-Fos expression, cellular growth factors come into focus (Sharpe et al., 1993; Ginty et al., 1994). Interestingly, p-CREB is involved in regulating BDNF transcription (Tao et al., 1998), which in turn binds to TrkB receptors, modulating synaptic long-term potentiation (Xu et al., 2000) and driving the expression of c-Fos and other early response genes.

Illustrating the richness of molecular changes due to manipulating sensory input, the expression of immediate early genes, among them egr-1 and c-fos, and neuronal plasticity-related genes such as those encoding for Arc, Syngr-1, and BDNF, was decreased by 2 weeks but increased again by 4 weeks in rat auditory cortex following bilateral cochlear ablation (Oh et al., 2007). ATF-2 is involved in the molecular underpinning of neuronal stability, cell death, and cellular growth (Yuan et al., 2009). Its expression goes down in retinal ganglion cells after their axotomy. Those cells capable of regrowing their axon show a return to high ATF-2 expression, indicating their return to normal conditions (Robinson GA, 1996). Members of the family of mitogen-activated protein kinases (MAPK) are ERK and p38 MAPK. ERK expression and phosphorylation reflects cellular changes associated to learning in an fear conditioning paradigm (Ota et al., 2010) and is co-regulated with c-Fos in several systems (Yang et al., 2008; Brami-Cherrier et al., 2009). Signals induced as a consequence of unilateral cochlear ablation are transduced mainly through the neuronal ERK pathway (Suneja & Potashner, 2003).

According to data available in the literature, the half-life of c-fos mRNA is 10-15 min (Müller et al., 1984; Sheng and Greenberg, 1990), whereas the c-Fos protein has a half-life of around 2 h (Curran et al., 1984; Müller et al., 1984).
1.3.3 Sensory stimulation of the auditory pathway

Sound-induced c-Fos expression has been employed to investigate the functional anatomy of the central auditory system of the mammalian brain. Expression of c-Fos has been seen in central auditory neurons after acoustical or electrical stimulation of the ear. Using pure-tone stimulation of mice or rats, the locations of neurons that turn positive for c-Fos or its mRNA were found to match the electrophysiologically established tonotopic maps in the ventral and dorsal cochlear nucleus (Rouiller et al., 1992; Brown & Liu, 1995; Miko et al., 2007), the superior olive (Adams, 1995), the dorsal nucleus of the lateral lemniscus (Saint Marie et al., 1999A), the inferior colliculus (Ehret & Fischer, 1991; Friauf, 1995; Pierson & Snyder-Keller, 1994; Saint Marie et al., 1999B), and the auditory cortex (Zuschratter et al., 1995). It was also induced in the vestibular nuclei (Sato et al., 1993). On exposing rats to specific behavioral tasks dependent on auditory stimuli, expression of c-Fos was also found in auditory centers of the diencephalon and telencephalon (Campeau & Watson, 1997; Carretta et al., 1999; Scheich & Zuschratter, 1995).

The amount of c-Fos mRNA also reflects habituation following stressful auditory stimulation (Campeau et al., 2002). Expression of c-Fos has also been used to estimate the driving force of axonal projections inside the central auditory system (Sun et al., 2009; Clarkson et al., 2010) and to localize the target site of audiogenic seizure (Kai & Niki, 2002) or the origin of tinnitus-related over-activity (Wu et al., 2003). The neuronal response reflected by c-Fos expression is related to the selective response of different subpopulations of neurons to sounds of time-varying properties (Lu et al., 2009) and may be induced by disinhibition following nerve lesions (Luo et al., 1999). The precision of a tonotopic c-Fos response to pure tone stimulation appears to be under the control of EphA4 and ephrin-B2 (Miko et al., 2007).

1.3.4 Electrical stimulation of the auditory pathway

Electrical intracochlear stimulation (EIS) as done with CIs in humans was shown to be similarly effective as acoustical stimulation in driving cells to express c-Fos (Rouiller et al., 1992; Illing & Michler, 2001; Nakamura et al., 2003). Subsequent studies showed a marked variability in the pattern of stimulation-dependent c-Fos expression in the auditory brainstem. Some studies reported massive induction of c-Fos in dorsal cochlear nucleus (DCN), with little expression in ventral cochlear nucleus (VCN), and a substantial increase of c-Fos immunoreactivity in the external nuclei of the inferior colliculus (IC), with little staining in its central nucleus CIC (Vischer et al., 1994; 1995; Zhang et al., 1996). Others reported distinctly different patterns of c-Fos expression upon EIS (Illing & Michler, 2001; Saint Marie et al., 1999A, 1999B; Saito et al., 1999; Zhang et al., 1998). In several studies involving electrical stimulation, tonotopic patterns of c-Fos expression were found (Saito et al., 1999; Nagase et al., 2000; Saito et al., 2000; Illing & Michler, 2001), but others did not report it (Nakamura et al., 2005). c-Fos has also been induced after neonatal auditory deprivation (Keilmann & Herdegen, 1995; 1997) and in kanamycin-deafened rats (Fujii et al., 1997; Nagase et al., 2003). Like EIS, directly stimulating the dorsal cochlear nucleus is effective in driving tonotopic c-Fos expression (Takagi et al., 2004) and produces hearing in rats (Zhang & Zhang, 2010).

1.4 GAP-43

In pre- and early postnatal development of the mammalian brain, expression of the membrane phosphoprotein GAP-43, also known as B-50, F1, pp46, P-57, or neuromodulin, is
high in neuronal somata, axons, and growth cones (Gispen et al., 1991; Kinney et al., 1993). Brain areas known for their adult potential for plasticity are characterized by high levels of GAP-43 (Benowitz et al., 1988; Benowitz and Routtenberg, 1997). This protein is a neuron-specific calmodulin-binding phosphoprotein and substrate for protein kinase C (Gispen et al., 1991; Schaechter & Benowitz, 1993). There are several lines of evidence relating this protein to axonal growth as well as to plasticity. It is produced at high levels in every nerve cell during neurite outgrowth and early stages of synaptogenesis (Skene and Willard, 1981; Mahalik et al., 1992) and represents a major constituent of the isolated growth cone (De Graan et al., 1985; Meiri et al., 1998). With maturation, its expression is down-regulated by most neurons (Skene, 1989; Benowitz & Perrone-Bizzozero, 1991). When a sense construct of GAP-43 mRNA was transiently expressed in non-neuronal cultured cells, these cells grow filopodial-like processes (Yankner et al., 1990; Verhaagen et al., 1994). If cells were transfected with a mutated construct of GAP-43 which prevented attachment of GAP-43 into the cell membrane, GAP-43 did not accumulate in pseudopods and no changes in cell morphology were induced (Widmer & Caroni, 1993). The attenuation of endogenous GAP-43 by an antibody that was raised against this protein and injected intracellularly has been found to reduce the degree of neurite outgrowth in a dose-dependent manner (Shea et al., 1992). The over-expression of GAP-43 in transgenic mice results in the formation of additional and aberrant neuronal connections (Aigner et al., 1995). Conversely, a knock out of the GAP-43 gene is survived by only 5-10% beyond weaning (Strittmatter et al., 1995). Whereas GAP-43 (-/-) mice show significant impairments in muscle strength, limb coordination and balance, and exhibit hyperactivity and reduced anxiety, GAP-43 (+/-) mice are only moderately impaired as compared with wild-type animals (Metz & Schwab, 2004). However, significant memory deficits were reported of heterozygous GAP-43 knockout mice with GAP-43 levels reduced by one-half, providing further evidence that GAP-43 exerts a crucial role in the bidirectional regulation of mnemonic processing (Rekart et al., 2005).

Expression of GAP-43 runs, at least partially, over activation of an AP-1 binding site. This identifies GAP-43 as a potential gene influenced by c-Fos. Further upstream, cooperation between p75(NTR) and TrkA results in an increased NGF-mediated TrkA autophosphorylation (Diolaiti et al., 2007). This cooperation also leads to a sustained activation of ERK-1/2 by phosphorylation and accelerates neurite outgrowth concomitant with a selective enhancement of the AP-1 activity and the transcriptional activation of genes such as GAP-43.

We discovered that unilateral deafness inflicted by a total sensory deafferentation of the cochlear nucleus in the mature rat invokes expression of GAP-43 (Illing & Horváth, 1995; Illing et al., 1997; see also Gil-Loyzaga et al., 2010). Since then, we found that this GAP-43 resides in presynaptic endings (Hildebrandt et al., 2011), is expressed by cholinergic neurons (Meidinger et al., 2006) residing in the ventral nucleus of the trapezoid body (Kraus & Illing, 2004), and innervate only specific subtypes of neurons in the cochlear nucleus (Illing et al., 2005). Altogether, these results clearly indicate that sensory deafferentation is followed by a specific reorganization of the neuronal network in the cochlear nucleus. Realizing that there is an impressive potential of neuroplasticity in the adult mammalian brainstem we wondered if other modifications of the pattern of sensory-evoked neuronal activity might be answered by a comparable response of the auditory brainstem in terms of synaptic reorganization and, as a consequence, signal representation and analysis. We therefore turned to EIS.
1.5 Linking c-Fos with GAP-43

Much is known about molecular networks regulating cellular growth and differentiation processes in general, but we still know little about the organization of the switch stands of activity-dependent molecular regulation in vivo. Regulatory molecules preceding c-Fos expression include p-CREB, p-ERK-1/2, p38 MAPK, and p-ATF-2. Molecules whose expression may be influenced by c-Fos include GAP-43, neuroskeletal elements, and cell adhesion molecules (Illing, 2001).

Williams et al. (1991) already suggested that an early response to sciatic nerve injury consisting of c-Fos expression in the spinal cord is transformed into a GAP-43 response emerging in the same spinal region of Rexed’s lamina II, if not the same cells. It remained open if axonal sprouting mediated by GAP-43 is beneficial to the lesioned systems or accounts for sensory disorders following nerve injury (Woolf et al., 1990).

Demonstrating a succession of c-Fos and GAP-43 in functional states of neurons related to activity-dependent modifications of brain structure more directly, Kleim et al. (1996) showed that the number of synapses per neuron and the percentage of Fos-positive cells within motor cortex was elevated in rats raised in conditions requiring acrobatic motor activity as compared to rats raised without such requirement. Their data suggest that Fos may be involved in the biochemical processes underlying skill acquisition and that motor learning, as opposed to motor activity, leads to increases in synapse number in the motor cortex. Correspondingly, Black et al. (1990) see a relation between c-Fos expression and synaptogenesis, and synaptogenesis, in turn, relies on the presence and action of GAP-43 (Benowitz & Routenberg, 1997). Diets improving spatial cognition of rats concomitantly raise the number of c-Fos positive neurons in hippocampus, resulting in a statistically significant negative correlation between the number of c-Fos positive neurons and the frequency of reference memory errors (Tanabe et al., 2004).

Watanabe et al. (1996) used a kindling model of epilepsy in which changes of neuronal activity in the form of brief focal seizures lead to lifelong structural and functional reorganization of the mammalian brain (McNamara et al., 1993). Brains thus affected by patterns of activity deviating from the normal develop axonal sprouting in hippocampal granule cells. However, when c-Fos was made unavailable for these brains due to a null-mutation, axonal sprouting is strongly reduced. Their data are consistent with the hypothesis that the lack of c-Fos reduces functional plasticity as well as a structural plasticity. This study again supports the hypothesis that the absence of c-Fos leads to a substantial attenuation of axonal plasticity due to a failure to activate growth-related genes such as GAP-43.

As a consequence of BDNF binding on the TrkB receptor, the expression of both c-Fos and GAP-43 is modulated (Edsjö et al., 2001; Koponen et al., 2004). Another feature shared by c-Fos mRNA and GAP-43 mRNA is to be stabilized by the same protein HuD (Chung et al., 1996; Mobarak et al., 2000; Smith et al., 2004). Binding of HuD apparently leads to increased levels of the respective protein under constant transcriptional conditions. This opens the possibility that c-Fos and GAP-43 share part of the regulation controlling their presence and availability, a feature that deserves attention in future research.
2. Methods

2.1 Animals

The region of the brain in which we looked for stimulation-dependent changes on the molecular and cellular level is the rat auditory brainstem. Wistar rats of either sex aged 6 to 12 weeks were used. Care and use of the animals as reported here were approved by the appropriate agency (Regierungspräsidium Freiburg, permission number 37/9185.81/G-10/83). Rats were anesthetized with a mixture of ketamin (i.p., 50 mg/kg body weight; Belapharm GmbH & Co. KG, Vechta, Germany) and xylazine (i.p., 5 mg/kg body weight; Rompun, Bayer-Leverkusen, Germany) before auditory brainstem response (ABR) measurements were done. For experiments involving acute EIS, anesthesia was induced by urethane (i.p., 1.5 g/kg body weight; Fluka AG, Buchs, Switzerland).

2.2 Deafness model

Between postnatal day (P) 10 and P20 (inclusive), neonatal Wistar rats received daily injections of kanamycin (i.p., 400 mg/kg body weight; Sigma, Taufkirchen, Germany) (Fig. 1). Due to hair cell destruction caused by this antibiotic (Matsuda et al., 1999; Osaka et al., 1979), a rise of hearing threshold was seen against normal hearing rats when rats have grown to adulthood (Fig. 2 A, B).

Fig. 1. Protocol to establish hearing and deaf experimental groups. Neonatal deafness was induced by daily kanamycin treatment between postnatal days 10 to 20. Measurement of acoustically or electrically evoked brainstem responses (ABR/EABR) were made around P33 and P42, respectively.

2.3 Electrical Intracochlear Stimulation (EIS)

Brains of hearing-experienced and hearing-inexperienced rats were analyzed after EIS lasting for different times, with stimulation durations spaced by the factor of 1.6: 45 min, 73 min, 2 h, 3:15 h, and 5 h (Fig. 2 C, D). The electrode used was a CI connected to a communicator both kindly provided by Cochlear GmbH (Hannover, Germany). Bipolar stimulation consisted of 50 Hz biphasic stimuli with a phase width of 50 µs. The electrically evoked brainstem response (EABR) was recorded like the ABR (see below) to determine an appropriate current level. The EABR was visualized using an averager (Multiliner E; Evolution 1.70c), calculating mean amplitudes over 500 sweeps in a frequency band of 0.1 to 10 kHz. We aimed to obtain maximal EABR amplitudes of 10 µV ± 10% by adjusting the current level of EIS to match acoustic stimuli of about 75 dB SPL. Chronic stimulation in the awake rat was done with the same mode using the same electrodes, but implants were connected to the stimulator by way of a swivel and a skull-based interface.
2.4 Auditory and electrically evoked brainstem responses

ABRs and EABRs were measured for hearing-experienced and hearing inexperienced rats. Hearing thresholds were tested acoustically (ABR) and electrically (EABR) by inducing auditory brainstem responses (Fig. 2 A, B). For ABR recording, steel needle electrodes were placed subcutaneously at vertex and mastoids and a 20 Hz train of click stimuli was presented to one side through a brass pipe equipped with a conical plastic tip into the ear canal, while sensations through the other ear were masked by supplying white noise at the same sound pressure level. Sound pressure was stepwise increased, attempting to elicit an ABR visualized by an averager (Multiliner E; Evolution 1.70c; Toennies, Germany). ABR mean amplitudes were determined after 300 sweeps in a frequency band of 0.1 to 3 kHz. If a kanamycin-treated rat showed an ABR at a sound pressure level smaller than 95 dB, recordings were stopped to avoid inducing further sensory stimulation. Following kanamycin treatment, we found ABR thresholds elevated by 75 dB to above 95 dB. These rats consistently failed to show a motor response to a handclap (Preyer's reflex). The absence of Preyer's reflex was taken by Jero et al. (2001) as indicating a rise of ABR threshold beyond 81 dB SPL. This result served as an additionally indication of deafness caused in the kanamycin-treated group.

![Fig. 2. ABR and EABR in hearing and deaf rats. Representative acoustically (A, B) and electrically (C, D) evoked auditory brainstem responses of hearing (A, C) and deaf (B, D) mature rats. Typically, the ABR of a hearing rat showed six distinguishable peaks (I-VI) and the EABR at least three peaks (II-IV). No ABR was detectable in kanamycin-treated rats up to 95 dB SPL. X-axis: 1 ms per unit; Y-axis: 4 µV per unit; EP: evoked potential.](www.intechopen.com)
immunocytochemistry based on diaminobenzidine staining or immunofluorescence (cp. Illing & Michler, 2001; Rosskothen et al., 2008; Rosskothen-Kuhl & Illing, 2010).

2.5 Immunohistochemistry (IHC)

After completion of the postoperative survival time or different stimulation periods, animals were killed by sodium-thiopental (i.p., 50 mg/ml per 200 g body weight of Trapanal 2.5 g, Nycomed, Konstanz, Germany) and perfused transcardially with a fixative containing 4% paraformaldehyde and 0% - 0.025% glutaraldehyde in 0.1 M phosphate buffer at pH 7.4. After brains were removed from the skulls and soaked in 30% sucrose overnight, parts containing anteroventral cochlear nucleus (AVCN), DCN, lateral superior olivary complex (LSO), and CIC were cryo-cut into 30 µm thick frontal sections. Following incubation with 0.05% H$_2$O$_2$, 1% sodium-borohydride (only for c-Fos and p-ATF-2) and 1% milk powder in 0.02 M phosphate buffer saline (PBS) at pH 7.4 for 30 min each, sections were exposed to a primary antibody either raised in goat against c-Fos (SC-52-G, 1:2000, lot. no. A2810, Santa Cruz Biotechnology Inc., Santa Cruz, USA), or raised in mouse against GAP-43 (MAB347, 1:5000, lot. no. LV1786431, Millipore, California, USA), or raised in rabbit against c-Jun p39 (SC-1694, 1:100–1:500; lot. no. L0606, Santa Cruz Biotechnology Inc.), p38 MAPK (4511S, 1:2000, lot. no. 5, Cell Signaling Technology, Inc., Danvers, USA), p-ATF-2 (5112S, 1:2000, lot. no. 10, Cell Signaling Technology, Inc.) or p-ERK 1/2 (SC-16982, 1:1000, lot. no. K1910, Santa Cruz Biotechnology Inc). After incubation for 48 h at 4° C, visualization of antibody-binding sites was based on DAB staining using biotinylated anti-goat/-mouse/-rabbit (BA-5000/BA2001/BA1000, 1:200, Vector Laboratories, Inc., Burlingame, USA) as secondary antibody and avidin-biotin-technique (Vector Laboratories) for signal intensification. Negative controls were run to verify specificity of the primary and secondary antibodies. Nuclei of the parabrachial region stained for c-Fos immunoreactivity served as positive controls (Illing et al., 2002).

2.6 In Situ Hybridization (ISH)

Thirty micrometer thick cryo-cut frontal brain sections were collected in 2x standard saline citrate (SSC) buffer (Invitrogen, Life Technologies GmbH, Darmstadt, Germany). The sections were washed in 2x SSC buffer for 15 min. Before pre-hybridization sections were pretreated in a 1:1 dilution of 2x SSC and hybridization buffer (50% formamide, Carl Roth GmbH, Karlsruhe, Germany), 4x SSC (Invitrogen), 10% dextran sulfate (Sigma, Taufkirchen, Germany), 1x Denhardt's solution (AMRESCO Inc., Ohio, USA), 250 µg/ml heat-denatured cod and herring sperm DNA (Roche Diagnostics GmbH, Mannheim, Germany), 625 µg/ml tRNA from E. coli MRE 600 (Roche) for 15 min. Pre-hybridization lasted in hybridization buffer for 60 min at 55°C. Hybridization was performed overnight at 55°C in the same solution with the addition of 100 ng/ml digoxigenin (DIG)-labeled c-Fos and 1000 ng/ml DIG-labeled GAP-43 antisense or sense cRNA, respectively. DIG-labeled sense and antisense cRNAs were synthesized by PCR amplification of brain tissue isolated c-Fos/GAP-43 mRNA. Primer design occurred by the use of the NCBI sequences NM_017195.3 (for GAP-43) and NM_022197.2 (for c-Fos). After hybridization, the sections were washed in 2x SSC for 2 x 15 minutes at room temperature, 2x SSC and 50% formamide (MERCK KGAA, Darmstadt, Germany) for 15 min, 0.1x SSC and 50% formamide for 15 min, and 0.1x SSC for 2 x 15 minutes at 65°C each. For immunological detection of DIG-labeled hybrids, brain
sections were treated in buffer 1 (100 mM Tris/HCl, pH 7.5) for 2 x 10 min each, blocked in buffer 2 (1% blocking reagent (Roche) in buffer 1) for 60 min at room temperature, and incubated overnight at 4°C with the anti-DIG Fab fragment from sheep tagged with alkaline phosphatase (1:1500, Roche) in buffer 2. For the color reaction, brain sections were equilibrated in buffer 1 for 2 x 10 min each and in buffer 3 (100 mM Tris/HCl, pH 9.5, 100 mM NaCl, 50 mM MgCl₂) for 10 min before the addition of nitroblue tetrazolium (0.34 mg/ml, Roche) and 5-bromo-4-chloro-3-indolyl-phosphate, 4-toluidine salt (0.17 mg/ml, Roche) diluted in buffer 3. Development of the color reaction was performed for around 9 hours in the dark at room temperature and stopped by transfer into aqua when the desired staining intensity was reached. Finally, sections were mounted on glass slides, dehydrated in increasing grades of alcohol, cleared in xylene, and coverslipped with DPX (Sigma).

2.7 Electron microscopy

After perfusion as described above, brains were postfixed in the same fixative for 2 h and stored overnight in PBS. Cutting was done on a microtome with vibrating blade (Leica VT 1000S, Bensheim, Germany) at 50 µm and collected in PBS containing 0.05 M glycine. Sections were permeabilized by a treatment with 5% and 10% dimethyl sulfoxide (DMSO), each for 10 min, and 20 min in 20% and 40% DMSO, all in PBS. Following washing in PBS, tissue was successively exposed to 0.05% H₂O₂, 0.05% sodium-borohydride, and 5% normal rabbit or goat serum in PBS each for 30 min, and incubated with antibodies raised against c-Fos (1:2000), p-ATF-2 (1:2000), or p-ERK-1/2 (1:1000). After incubation for 24 h at 4°C, visualization of antibody-binding sites occurred as described before for the IHC protocol (see. 2.5). Afterward, brain sections were incubated with 0.1% osmium-tetroxide in 0.1 M cacodylic acid buffer (pH 7.4), dehydrated in ethanol and embedded in EMbed-812 (Science Services, München, Germany).

3. Results and discussion

3.1 Effectiveness of EIS to induce gene expression

EIS is an effective way to activate neuronal networks in the central auditory system, including the induction of c-Fos expression throughout the brainstem of hearing-experienced rats (Fig. 3).

We monitored c-Fos expression on the transcriptional level by detecting c-Fos mRNA as well as on the translational level by c-Fos immunoreactivity (Illing et al., 1999). In both cases we detected staining in neuronal nuclei that closely matched in space and time (Fig. 4).

Attempting to understand the laws according to which neurons express c-Fos as a consequence of EIS we, first, identified the cell types in which c-Fos emerged. Following 2 h of EIS we found that only distinct subpopulations of neurons in VCN, DCN, LSO, and CIC express c-Fos (Reisch et al., 2007) (Fig.5). Whereas sub-populations of glutamatergic and glycinergic cells responded in all four regions, GABAergic neurons failed to do so except in marginal zones of AVCN and in DCN (but cp. Ishida et al., 2002). Combining immunocytochemistry with axonal tracing, neurons participating in major ascending pathways, commissural cells of VCN and certain types of neurons of the descending auditory system were seen to respond to EIS with early response gene expression. By contrast, principal LSO cells projecting to the contralateral CIC as well as collicular efferents of the DCN did not.
Fig. 3. Effectiveness of EIS to induce c-Fos expression throughout the brainstem of hearing rats. (A) Tonotopic c-Fos expression (black dots) in AVCNi after 2 h of EIS. (B) Tonotopic band (arrowheads) of c-Fos positive nuclei (black dots) in CICc after 3:15 h of EIS. (C) Tonotopic c-Fos expression (black dots) in the deep layers of DCNi after 5 h of EIS. (D) c-Fos expression (black dots) in LSOi following 5 h of sustained EIS. The dashed lines correspond to the borders of the respective auditory regions. Scale bars: 0.2 mm. i: ipsilateral; c: contralateral; CB: cerebellum; n7: facial nerve.

Fig. 4. Comparison of c-Fos immunoreactivity (A, C) vs. c-Fos in situ hybridization (B, D) in AVCNi of a hearing-experienced rat after 73 min EIS. (A, B) The pattern of c-Fos protein (black dots) and c-Fos mRNA (blue dots) of 2 adjacent sections is nearly identical. Scale bar: 0.2 mm. (C, D) Higher magnification of c-Fos protein positive nuclei (C) and c-Fos mRNA positive neurons (D) in AVCNi. Scale bars: 20 µm. i: ipsilateral; d: dorsal; l: lateral.
Fig. 5. EIS-dependent c-Fos expression in neurons of the auditory brainstem of hearing-experienced rats (Illing et al., 2010). (A) In VCN, half of the c-Fos positive nuclei were localized in glycinergic cells (Gly, yellow, asterisk), the other half showed no double labeling with glycine (arrow). (B) c-Fos in a large VCN commissural neuron after injection of the tracer Fast Blue into VCNc. (C) c-Fos in a GABAergic neuron of DCN. (D) c-Fos positive nuclei were never observed in neurons of DCN projecting to the contralateral CICc. (E) Some (arrow) but not all (asterisk) lateral olivocochlear (LOC) neurons labeled from the cochlea by axonal tracing showed co-localization with c-Fos. (F) c-Fos positive nuclei in small glutamatergic cells of LSOi (Glu, arrow), large glutamatergic cells lacked c-Fos immunoreactivity (asterisk). (G) CIC lacking c-Fos positive GABAergic cells. (H) Immunoreactivity for Egr-1 coincided with c-Fos immunoreactivity. i: ipsilateral, c: contralateral. Scale bars: 20 µm in D, 10 µm for all others.

Second, following continuous stimulation of hearing-experienced anesthetized rats for various durations from 45 min to 5 hours, we observed a non-linear increase of the population of c-Fos positive neurons, with a rise followed by a decline, followed by a second rise of the number of stained cells in VCN (Rosskothen-Kuhl & Illing, 2010) (Fig. 6).

Fig. 6. Pattern of c-Fos expression with increasing stimulation time in AVCNi of hearing-experienced rats. (A) In control rats, AVCNi is devoid of c-Fos staining. (B-D) Typically, the number of c-Fos positive cells (black dots) increased from 45 min to 2 h in a region tonotopically corresponding to the stimulation position. (E-F) By 3:15 h of unilateral EIS, the c-Fos level transiently decreases before it rose again by 5 h (F). Scale bar: 0.2 mm. Co: control; i: ipsilateral; CB: cerebellum; d: dorsal; l: lateral.
Third, we used different temporal patterns known to have major significance for central auditory processing (Walton, 2010) for EIS at a constant duration of 2 h (Jakob & Illing, 2008). For each of the major auditory brainstem nuclei and some of their subregions we found specific patterns of the immunoresponsive cell populations reflecting different aspects of the stimulation site and stimulation parameters (Table 1).

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Table 1. Expression pattern of c-Fos across the auditory brainstem depends on stimulation parameters. Each of the major auditory brainstem regions, and some of their subregions, showed a unique fingerprint of c-Fos expression (cp. rows) with respect to reflecting laterality, intensity, frequency (Jakob & Illing, 2008), and duration of EIS. MGB: medial geniculate nucleus, i: ipsilateral, c: contralateral.

### 3.2 Pharmacology

Apart from the duration of EIS and the particular stimulation parameters used, the possibility exists that the consequences of activating the central auditory system may also be

![Fig. 7](https://www.intechopen.com)

Fig. 7. Effect of Nimodipine on the pattern of c-Fos expression in the DCNi with different stimulation parameters. (B-D, F-H) For both experimental groups a shift of c-Fos positive nuclei (black dots) from the upper to the deep layers of DCN exist with increasing stimulation frequency, after 2 h of EIS. However, the Nimodipine treatment (E-H) resulted in a significant increase in number of c-Fos expressing neurons compared to non-treated (A-D) rats. Scale bar: 0.2 mm. Co: control; i: ipsilateral; d: dorsal; l: lateral.
influenced by pharmacological manipulations. Here we illustrate one example, the impact of Nimodipine systemically administered before EIS. In DCN, the pattern of neurons expressing c-Fos is dramatically different depending on the presence or absence of Nimodipine (Fig. 7).

3.3 Molecules upstream of c-Fos expression

In order to understand the functional relevance of c-Fos expression under any particular mode of stimulation, it is mandatory to co-localize it with other molecular actors of neuroplastic change in space and time. Therefore, we studied the spatio-temporal relationship of c-Fos expression with the expression pattern of molecules known to be involved directly or indirectly in the regulation of c-Fos expression. Upstream of c-Fos expression, p-CREB, ATF-2, p-ERK-1/2, and p38 MAPK unfold their activity. We discovered that their expression is modulated either positively or negatively under EIS (Fig. 8).

Fig. 8. Modulation of regulatory molecules upstream of c-Fos expression. (A-B) 2 h after EIS of hearing rats p-CREB positive neurons were rare in a middle band of AVCNi (A, arrow) corresponding to the c-Fos positive area in the adjacent section (B, arrow). (C-D) A similar pattern existed for p-ATF-2 (C, arrowheads) and c-Fos (D, arrowheads) positive nuclei in the AVCNi of deaf rats after 2 h of EIS. (E-H) 2 h after EIS of deaf rats p-ERK-1/2 immunoreactivity was detected only in the LSO of the stimulated side (E, arrows) corresponding to the c-Fos expression pattern (not shown). Like for p-ERK-1/2 the p38 MAPK immunoreactivity was increased in the medial LSOi (G, arrows) compared to the contralateral side (H). c: contralateral; i: ipsilateral. Scale bar: 0.2 mm.

Next, we looked for the ultrastructural localizations of molecules involved in the stimulation-dependent regulation of transcription factors (Fig. 9). Under EIS, c-Fos emerged in nuclei and rough endoplasmic reticulum of VCN neurons. p-ATF-2 and p-ERK-1/2 showed specific regional and intracellular staining patterns of their own.

Directly interacting with c-Fos is c-Jun p39. We found that, similar to the immediate early gene egr-1 (De et al., 2003) but unlike its stimulation-dependent expression in the inner ear (Ruan et al., 2007), c-Jun is present in most of the auditory brainstem in normal brains without specific stimulation (Fig. 10). Upon EIS and ensuing c-Fos expression, c-Fos and c-Jun may dimerize to form AP-1 in a highly active variant (Rosskothen et al., 2008) (Fig. 10 B-D).
Fig. 9. Modulation of transcription factor regulators by EIS. (A) Ultrastructural localization of c-Fos immunoreactivity in nucleus (asterisk) and cytoplasm (arrows) of a neuron in AVCNi. (B) Contralaterally, no such staining was seen. (C) p-ATF-2 immunoreactivity in the nucleus of an AVCNi neuron; upon EIS, this staining declines. (D) p-ERK-1/2 staining in nucleus (asterisk) and cytoplasm (arrows) in a DCN neuron next to an unstained neuron after EIS. Dashed lines indicate cell boundaries. c: contralateral; i: ipsilateral. Scale bar: 2 µm.

3.4 Molecules downstream of c-Fos expression

Downstream of c-Fos lie many genes regulated by an AP-1 binding site in their promoter, among them the gene encoding GAP-43. We observed an emergence of GAP-43 in fibers and boutons of VCN on the side of chronic EIS (Fig. 11). At the same time, GAP-43 mRNA is strongly up-regulated in neurons of LSO (Fig. 11 C-F).

The LSO is known to be involved in relearning sound localization during unilateral conductive hearing loss (Irving et al., 2011), suggesting that its neurons receive a growth signal by unilateral EIS that, in turn, causes axonal sprouting in VCN on the stimulated side. This scenario poses a remarkable contrast to GAP-43 expression in VCN one week after its total sensory deafferentation by cochlear ablation (Illing et al., 1997). Under these circumstances, GAP-43 in VCN is supplied by neurons of the ventral nucleus of the trapezoid body (VNTB), source of the medial olivocochlear pathway, rather than of LSO, source of the lateral olivocochlear pathway.
3.5 EIS induces c-Fos expression in deaf rats

The absence of hearing experience has far-reaching consequences for the interneuronal communication within networks of the auditory brainstem. First, when hearing fails, EIS
entails expression of c-Fos in populations of neurons that are much larger than normally, essentially disregard tonotopic order, and lack much of spatio-temporal variations as seen in hearing-experienced rats (Rosskothen-Kuhl & Illing, 2012) (Fig. 12).

Fig. 12. Differences of c-Fos expression pattern in hearing vs. deafened rats. (A) Tonotopic c-Fos expression (black dots) in AVCNi of a hearing rat after 2 h of EIS. (B) Tonotopic band (arrowheads) of c-Fos positive nuclei (black dots) in CICc of a hearing rat after 3:15 h of EIS. (C) Non-tonotopic c-Fos expression (black dots) in AVCNi of a deaf rat after 2 h of EIS. (D) Strong and extended c-Fos expression (black dots) in the dorsolateral part of the CICc of a deaf rat after 3:15 h of EIS. i: ipsilateral; c: contralateral; CB: cerebellum; CX: Neocortex. Scale bars: 0.2 mm.

Second, the composition of c-Fos expressing subpopulation changed with the preceding hearing experience (Fig. 13).
Fig. 13. Composition of c-Fos expressing subpopulations changed depending on preceding hearing experience and stimulation time. (A) Co-localization (arrow) of c-Fos positive nuclei (red, asterisk) and parvalbumin (PV)-positive neurons (green, arrowhead) in the AVCNi after EIS. (B, C) Statistic evaluation of co-localization of c-Fos positive nuclei and PV-positive neurons resulted only for hearing rats (B, p=0.018) but not for deaf rats (C) in a significant increase of double-labeling from 2 to 5 h of stimulation. (D) Following EIS, c-Fos expression (green, asterisk) was also co-localized (arrow) with GABAergic somata (red, arrowhead) in marginal zones of AVCNi. Scale bar: 20 µm. (E, F) The statistic evaluation of co-localization of c-Fos positive nuclei and GABA-positive neurons resulted only for hearing rats (E, p=0.006) and not for deaf rats (F) in a significant increase of double-labeling from 2 to 5 h of EIS. i: ipsilateral; ns: non-significant.

4. Conclusions and outlook

The brain is responsive to changes of activity in sensory nerves by forming memory traces on all its levels, from forebrain to hindbrain, and from molecules to neuronal networks. A successful therapy for patients with dysfunctional ears requires a CI that plays patterns of electrical activity corresponding to patterns of sound to the auditory nerve, and a moldable brain responding to this pattern. This response must include the initiation of molecular and synaptic reorganization to make optimal use of sensory-evoked activity, specifically to identify and categorize temporal patterns in spoken language (Munro, 2008; Anderson et al., 2010; Skoe & Kraus 2010). Obviously, then, it is fundamentally important to run the CI in a way that fully exploits the neuroplastic potential of the central auditory system. It is therefore essential to detect short and long term molecular and cellular changes in the central nervous system in response to EIS equivalent to CI stimulation to understand which stimulation parameters are more important than others to elicit a full and focused plasticity response.

As we showed, specific conditions of afferent activity prompt specific populations of neurons in specific regions of the auditory brainstem to prepare for a molecular, structural,
and functional remodeling within hours. Laterality, frequency, duration, and intensity of the stimulus each affect different neuronal populations across different regions. Coding of laterality takes place below the midbrain, but appears to be integrated in network activity from the inferior colliculus upwards. Each major brainstem region involved in the analysis of auditory stimuli respond to specific stimulation parameters by a unique dynamic pattern of c-Fos expression. With our studies we have made a first and perhaps a second and third step to understand central auditory plasticity, and there is reason to consider much of our data obtained from a mammalian brain transferable to man. However, our studies are not yet comprehensive in the sense that they already provide binding recommendations to CI programmers or users. Instead, it provides data showing for the first time that the parameters of running a CI bear a very real potential for neuroplastic remodeling of the central auditory system, and that the brain is certain to show a wealth of molecular and cellular responses to CI stimulation.

More stimulation parameters must be tested and more molecular markers monitored in order to delineate robust relationships between modes of stimulation, molecular processes, cellular growth, network remodeling, and the efficiency of signal processing. The knowledge so obtained will set the stage for the development of CIs in the future.

5. Acknowledgements

It is a pleasure to express our gratefulness to many co-workers for help: Heika Hildebrandt, Ann-Kathrin Rauch, Peter Bischoff, Till Jakob, Ulrike Doering, Michaela Fredrich, Eike Michalk, Peter Pedersen, Jürgen John, Alexander Huber, Ralf Birkenhäger, and Sigrid Weis. Moreover, thanks go to Cochlear Deutschland GmbH for kindly providing hard- and software used for EIS, and to Roland Laszig for continuous support.

6. List of abbreviations

2DG, 2-desoxy-D-glucose
ABR, auditory brainstem response
AP-1, activator protein-1
Arc, activity-regulated cytoskeleton-associated protein
ATF-2, activating transcription factor-2
AVCN, anteroventral cochlear nucleus
AVCNc, AVCN contralateral to stimulation
AVCNi, AVCN ipsilateral to stimulation
B50, synonym for GAP-43
BDNF, brain derived neurotrophic factor
bFGF, basic fibroblast growth factor, same as FGF2
c, contralateral
CaMKIV, Ca\(^{2+}\)/calmodulin-dependent protein kinases IV
cAMP, cyclic adenosine monophosphate
CB, cerebellum
c-fos, Finkel-Biskin-Jinkins murine osteosarcoma viral oncogene
c-Fos, protein encoded by proto-oncogene c-fos
CI, cochlear implant
CIC, central nucleus of the inferior colliculus
CICc, CIC contralateral to stimulation
c\textit{jun}, avian sarcoma virus 17, 'Jun' being derived from Japanese 'ju-nana' for '17'
c-Jun, protein encoded by immediate-early gene c\textit{jun}
Co, control
CRE, cAMP response element
CREB, cAMP response element-binding protein
c\textit{RNA}, coding ribonucleic acid
CX, neocortex
d, dorsal
DAB, 3,3'\text{\textprime}diaminobenzidine tetrahydrochloride
dB, decibel
DCN, dorsal cochlear nucleus
DCNi, DCN ipsilateral to stimulation
DIG, digoxigenin
DMSO, dimethyl sulphoxide
DNA, deoxyribonucleic acid
EABR, electrical auditory brainstem response
Egr-1, early growth response protein-1
EIS, electrical intracochlear stimulation
EP, evoked potential
EphA4, ephedrine A4
ERK, extracellular signal regulated kinase, mitogen-activated protein (MAP) kinases, variants 1 and 2
F1, synonym for GAP-43
FAP, activator protein-1 (AP-1)-like sequence
GABA, gamma amino butyric acid
GAP-43, growth-associated protein-43
Glu, glutamate
Gly, glycine
HuD, member of human Hu proteins identified as target antigens in autoimmune paraneoplastic encephalomyelitis-sensory neuronopathy, RNA-binding and stabilizing protein.
Hz, Hertz
i, ipsilateral
i.p., intraperitoneal
IC, inferior colliculus
IHC, immunhistochemistry
ISH, in situ hybridization
Krox-24, synonym for Egr-1
l, lateral
LOC, lateral olivocochlear
LSO, lateral superior olivary complex
LSOc, LSO contralateral to stimulation
LSOi, LSO ipsilateral to stimulation
m, medial
MAPK, mitogen-activated protein kinase
MGB, medial geniculate body
MGBc, MGB contralateral to stimulation
mRNA, messenger ribonucleic acid
n7, facial nerve
NGF, nerve growth factor
NGFI-A, nerve growth factor-induced protein 1, synonym for Egr-1
NMDA, N-methyl-D-aspartic acid
ns, non-significant
P, postnatal day
p38, protein phosphorylating specific MAPKs
P-57, synonym for GAP-43
p75(NTR), p75 neurotrophin receptor
p-ATF-2, phosphorylated activating transcription factor-2
PBS, phosphate-buffered saline
PCR, polymerase chain reaction
p-CREB, phosphorylated cAMP response element binding protein
p-ERK, phosphorylated ERK
pp46, synonym for GAP-43
PV, parvalbumin
SIE, sis inducible element associated to retroviral DNA sequences (v-sis) originally isolated from simian sarcoma virus
SPL, sound pressure level
SRE, serum response element
SSC, standard saline citrate
Syngr-1, Synaptogyrin-1
Tis8, synonym for Egr-1
TrkA, neurotrophic tyrosine kinase receptor type 1
TrkB, neurotrophic tyrosine kinase receptor type 2
tRNA, transfer ribonucleic acid
VCN, ventral cochlear nucleus
VCNc, VCN contralateral to stimulation
VNTB, ventral nucleus of the trapezoid body
Zenk, synonym for Egr-1
Zif268, synonym for Egr-1

7. References


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For many years or decades, cochlear implants have been an exciting research area covering multiple disciplines which include surgery, engineering, audiology, speech language pathology, education and psychology, among others. Through these research studies, we have started to learn or have better understanding on various aspects of cochlear implant surgery and what follows after the surgery, the implant technology and other related aspects of cochlear implantation. Some are much better than the others but nevertheless, many are yet to be learnt. This book is intended to fill up some gaps in cochlear implant research studies. The compilation of the studies cover a fairly wide range of topics including surgical issues, some basic auditory research, and work to improve the speech or sound processing strategies, some ethical issues in language development and cochlear implantation in cases with auditory neuropathy spectrum disorder. The book is meant for postgraduate students, researchers and clinicians in the field to get some updates in their respective areas.

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