

REVIEW

Building neuromuscular junctions in vitro

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ABSTRACT

The neuromuscular junction (NMJ) has been the model of choice to understand the principles of communication at chemical synapses. Following groundbreaking experiments carried out over 60 years ago, many studies have focused on the molecular mechanisms underlying the development and physiology of these synapses. This Review summarizes the progress made to date towards obtaining faithful models of NMJs *in vitro*. We provide a historical approach discussing initial experiments investigating NMJ development and function from *Xenopus* to mice, the creation of chimeric co-cultures, *in vivo* approaches and co-culture methods from *ex vivo* and *in vitro* derived cells, as well as the most recent developments to generate human NMJs. We discuss the benefits of these techniques and the challenges to be addressed in the future for promoting our understanding of development and human disease.

KEY WORDS: Neuromuscular junctions, *In vitro* models, Human pluripotent stem cells

Introduction

At the beginning of the last century the 'neuron doctrine' proposed by Ramón y Cajal revolutionized the emerging field of neuroscience. The doctrine firmly established individual neurons as the basic anatomical and physiological unit of the nervous system, and prompted many questions regarding how neurons communicate with one another and how the space between them is crossed by the signals responsible for that communication. We now know that neurons make complex structures called synapses that allow chemical transmission of signals among cells. The neuromuscular junction (NMJ) is a tripartite chemical synapse of the peripheral nervous system. It allows the transmission of motor information that governs muscle contraction from the central nervous system to the muscle fibers. The NMJ comprises a motoneuron on the presynaptic side, a muscle cell on the postsynaptic side and a terminal Schwann cell (tSC) wrapping the NMJ structure. A fourth cell type, a kranocyte, which caps the NMJ and plays a role in synapse regeneration, has recently been identified in rodents (Court et al., 2008) (Fig. 1). Fatt and Katz showed that endplate potentials (EPPs), which correspond to the chemically induced change in electric potential of the motor end plate, originate from the spontaneous or electrically evoked release of acetylcholine (ACh) from the nerves (Fatt and Katz, 1951). Furthermore, Katz and colleagues proposed the 'quantal hypothesis', stating that ACh is released from the nerve terminal in packets or 'quanta'. The spontaneous release of one quantum results in a miniature EPP (mEPP) in the post-synaptic cell, whereas the synchronized release

at discrete zones of the nerve terminal called 'active zones' (Couteaux and Pecot-Dechavassine, 1970). A functional definition of active zones came a few years later, when it was shown that vesicles dock and fuse at the membrane, and their contents are exocytosed in response to nerve depolarization (Heuser et al., 1979). Taken together, these pioneering studies on NMJs provided the basis for our current knowledge of the intricate mechanisms underlying synaptic transmission.

Neuromuscular junctions have since been the most intensively

of multiple ACh quanta leads to EPPs. Synaptic vesicles accumulate

studied synapses, because of their comparatively large size, relative simplicity and accessibility. Indeed, among specific features, neurotransmitter release at this synapse is mediated by multiple docking sites present in long active zones in frogs or several active zones in mammals. In contrast, synaptic transmission in the brain is associated with a single active zone per synapse (Biro et al., 2005; Silver et al., 2003). Another important distinction between NMJs and central synapses is that the post-synaptic element at an NMJ only receives cholinergic inputs, whereas post-synaptic neurons receive a large variety of inputs, which makes the analysis of the response to a single neurotransmitter more difficult to interpret. Therefore, the NMJ remains an excellent model for studying the mechanisms of neurotransmission, and is well suited to study synaptic formation, maintenance and function. Furthermore, the NMJ is at the core of a number of human pathologies that all have the perturbation of neurotransmission in common (Box 1).

Here, we provide an update on the different *in vitro* models that have been investigated to increase our understanding of the normal and pathological development of NMJ. We present a focus on the recent findings on humanized *in vitro* models, which have been made possible using human pluripotent stem cells, and discuss the promises and challenges of these new models.

The neuromuscular junction: a model system for studying synaptic development and function

A great deal of information on how synapses form has been obtained through studying muscle cell lines (mostly the murine C2C12 cell line) that form spontaneous acetylcholine receptor (AChR) aggregates, and co-cultures of muscle cells and motoneurons (Sanes and Lichtman, 2001). Early studies were carried out in cultures of neurons and muscles from Xenopus laevis (Anderson et al., 1977) or from chick (Frank and Fischbach, 1979), laying the groundwork for understanding the mechanisms of in vitro NMJ formation and the study of NMJ disorders (discussed below). These datasets have been extended in vivo with the emergence of transgenic mice, along with the use of nonmammalian animal models (Box 2). Here, we briefly describe the steps in NMJ formation as well as the essential molecular and structural markers of NMJ formation and maturation (Figs 1 and 2). For more detailed discussion of the mechanisms of synapse formation, we refer the reader to recent reviews (Burden et al., 2018; Li et al., 2018).

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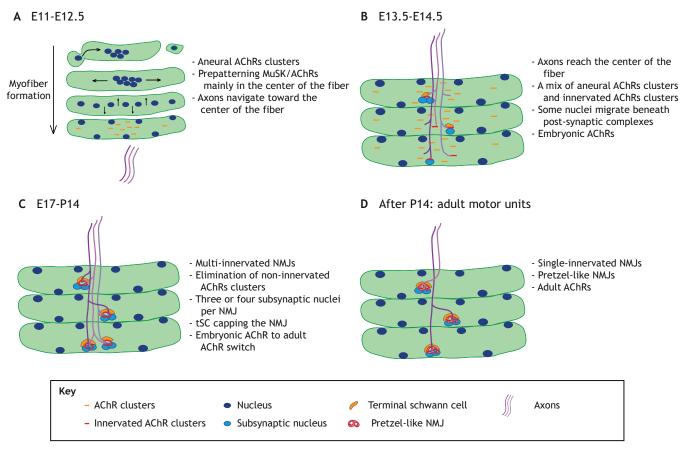


Fig. 1. The stages of NMJ formation in mice. (A) During development, prior to and independent of innervation, AChRs are clustered in the prospective region of muscle. Thus, the muscle cell plays an instructive role from a very early stage of development, in positioning and restricting synapses to a central band in the muscle, and building a post-synaptic density prior to axonal contact. (B) Axons do not contact muscle fibers at random locations, but rather in a narrow central region of the muscle fiber characterized by the presence of aneural 'pre-patterned' AChR clusters (Lin et al., 2001; Yang et al., 2001). Axons then become apposed to AChR clusters. (C) Additionally three or four nuclei accumulate under the post-synaptic membrane in the muscle cells (which are defined as multinucleated syncytia formed during development by fusion of mononucleated precursor cells) and become specialized for transcription of synaptic proteins. Concomitantly, folds form in the muscle membrane increasing the area of contact between the post-synaptic membrane and the extracellular matrix (Guarino et al., 2019). (D) Muscle fibers are poly-innervated until the end of the second week after birth in rodents. This is when the multiple innervations of individual post-synaptic densities are reduced to a single innervating motor axon through a process called synapse elimination (Sanes and Lichtman, 2001). AChR, acetylcholine receptor; adult AChR, $\alpha 2\beta \epsilon \delta$; E, embryonic day; embryonic AChR, $\alpha 2\beta \gamma \delta$; MuSK, muscle-specific kinase; P, post-natal day; tSC, terminal Schwann cell.

Cellular components in synapse formation

NMJ development requires extensive communication among the three key components of the tripartite synapse: presynaptic motoneurons, post-synaptic muscle fibers and tSCs. NMJ formation is closely intertwined with myogenesis and innervation occurs between embryonic days (E) 12.5 and E14 in mice, immediately after the first wave of myotube formation. This timing varies slightly among specific muscles and their locations (Fig. 1).

The composition of AChRs changes with development as follows: nicotinic AChRs are pentamers composed of five subunits in a stoichiometry of two α -, one β -, one δ - and one γ -subunit in the embryo. These are replaced by a ϵ -subunit at the postnatal age (Changeux et al., 1992). Within 3 weeks of birth, AChR clusters adopt a complex, pretzel-like configuration in mice (Sanes and Lichtman, 2001). The maturation of the NMJ leads to an increase in number, stability, density and clustering of AChRs in the post-synaptic compartment, as well as the differentiation of the presynaptic membrane into active zones with proper synaptic proteins and associated synaptic vesicles (Burden et al., 2018). The overall process reflects strong reciprocal signaling between pre- and postsynaptic structures.

Another crucial component involved in the formation of the NMJ is the tSC. Although the very first steps in synapse formation do not require the presence of tSCs, these cells are necessary for maintenance and maturation of NMJs (Ko and Robitaille, 2015). Ablation of tSCs after NMJ maturation at P30 in mice, leads to NMJ fragmentation and neuromuscular transmission deficits (Barik et al., 2016).

Molecular components of NMJ development and function

Over the past decade, extensive progress has been made in understanding the molecular mechanisms involved in NMJ formation. Here, we summarize the current knowledge on some of the key molecules and signaling pathways that are indispensable for NMJ formation and maintenance (Figs 1 and 2).

MuSK-Lrp4-agrin signaling

MuSK (muscle-specific kinase), a transmembrane receptor tyrosine kinase and its co-receptor LRP4 (low-density lipoprotein receptor-related protein 4) are crucial for the aggregation of AChRs before and after innervation. Prior to innervation, the MuSK/LRP4 complex accumulates in the muscle central region where it induces AChR

Box 1. Neuromuscular junction disorders

Neuromuscular diseases (NMD) refer to a vast group of conditions that perturb or even progressively block the control of muscles. NMDs can originate from motoneuron defects, e.g. as in amyotrophic lateral sclerosis (ALS) and in spinal muscular atrophy (SMA), or from a defect of the neuromuscular junction itself, e.g. as in congenital myasthenic syndromes (CMS), and in autoimmune diseases such as myasthenia gravis (MG). NMJ diseases have a wide range of origins, including genetic mutations, antibodies or even toxins that interfere with the control of signaling between the presynaptic nerve ending and the post-synaptic muscle membrane, and/or the formation of the synapse.

Myasthenia gravis (MG) is one of the most common diseases of the NMJ. This autoimmune disorder of the neuromuscular transmission is increasingly considered as a syndrome more than as a single disease. At least three different antibodies to AChRs, MuSK or LRP4 have been identified as the cause of the typical weakness and fatigability associated with MG (Higuchi et al., 2011; Hoch et al., 2001; Lindstrom et al., 1976). In addition, Lambert-Eaton myasthenic syndrome (LEMS) is caused by auto-antibodies against voltage-gated calcium channels (Lennon et al., 1995).

As well as these autoimmune diseases, there is also a large family of genetic conditions that affect the functioning of the NMJ. Remarkable progress in molecular genetics during recent decades has allowed the identification of an increasing number of mutations in genes encoding proteins of the NMJ that drive pathogenesis of several inherited diseases. Mutations in key human genes, such as *DOKT*, *RAPSN*, *MUSK*, *LAMB2*, *COLQ*, *AGRN* and genes encoding AChRs, can cause different forms of CMS (Beeson et al., 2006; Engel et al., 2015; Hantai et al., 2013; Ohkawara et al., 2014; Verschuuren et al., 2016).

clustering. With innervation, LRP4 binds a neural isoform of agrin (AGRN), a heparan sulfate proteoglycan secreted by the axon terminal (Ferns et al., 1993; Kim et al., 2008; Ruegg and Bixby, 1998; Zhang et al., 2008). Agrin interaction with LRP4 results in the formation of tetrameric agrin-LRP4 complexes that promote activation and transphosphorylation of MuSK, leading to AChR clustering and NMJ formation (Zong et al., 2012). AChR clusters that are apposed to nerve terminals persist and grow, while the non-apposed ones disassemble (Wallace, 1989). Mutations in the human *AGRN* gene cause congenital myasthenic syndrome (CMS), with distal muscle weakness and atrophy (Nicole et al., 2014).

More recently, two studies have independently revealed that the muscle LRP4 also acts as a retrograde signal for presynaptic differentiation (Wu et al., 2012; Yumoto et al., 2012). MuSK activation requires Dok7 (docking protein 7; an adaptor protein expressed by the muscle), which is indispensable for MuSK activity and therefore AChR clustering. The binding of Dok7 to the cytoplasmic juxtamembrane region of MuSK strongly enhances the tyrosine kinase activity of MuSK, leading to its activation (Hamuro et al., 2008; Inoue et al., 2009). MuSK-deficient mice show neither muscle pre-patterning of AChRs nor NMJ formation, and die perinatally from respiratory failure (Okada et al., 2006). More recently, it has been shown that the postnatal knockdown of *Dok7* gene in mice causes structural defects in NMJs and motor dysfunction, supporting a role for Dok7 in NMJ maintenance (Eguchi et al., 2016).

AChR-rapsyn

Beginning around the time of birth, fetal AChRs ($\alpha 2\beta\gamma\delta$) are gradually replaced by the adult subtype of AChRs ($\alpha 2\beta\epsilon\delta$). Although the physiological significance of this developmental switch remains to be fully understood, different studies have suggested a role of the γ -subunit during the NMJ formation and

more specifically in innervation patterning (Koenen et al., 2005; Takahashi et al., 2002). For example, deletion of the mouse AChR γ-subunit leads to perinatal lethality associated with an absence of pre-patterned AChR clusters during initial stages of neuromuscular synaptogenesis (Koenen et al., 2005; Takahashi et al., 2002). A key effector in AChR clustering is the protein rapsyn, a cytoplasmic scaffolding protein expressed in the muscle, which binds directly to AChR and anchors AChR in the post-synaptic membrane (LaRochelle et al., 1990; Sobel et al., 1978). Rapsyn-deficient mice fail to develop post-synaptic specializations (Gautam et al., 1995), indicating that rapsyn is essential for the tethering of AChRs to the post-synaptic apparatus. Although rapsyn assembles with AChRs in the secretory pathway (Marchand et al., 2002; Moransard et al., 2003), the exact mechanisms downstream of MuSK/Dok7 that stimulate rapsyn-induced AChR clustering remain to be determined.

Extracellular components

In addition to these core molecules that are essential for NMJ formation, a number of important regulators also participate in this process. Whis are a group of glycoproteins secreted at the NMJ by the muscle fiber, motoneurons and tSCs. Several Whit ligands bind to the cysteine-rich domain (CRD) in MuSK (Barik et al., 2016), and play a role in axon extension and branching, as well as in AChR prepatterning (Li et al., 2018). Both canonical and non-canonical planar cell polarity (PCP) Whit pathways are affected in transgenic mice in which the CRD in MuSK is deleted (Messeant et al., 2017).

Dystroglycan (DG) has a dual function: it organizes extracellular matrix (ECM) scaffolds and stabilizes AChR clusters. DG is formed of two subunits that are generated by proteolytic cleavage of a single precursor protein. The extracellular subunit is necessary to organize a functional scaffold in the basal lamina including perlecan, acetylcholinesterase/ColQ and laminin (Jacobson et al., 2001; Peng

Box 2. Contributions of non-mammalian model organisms for NMJ understanding

In parallel to the *in vitro* models for NMJ described in this Review, non-mammalian model organisms such as *Drosophila* and zebrafish have also been instrumental in identifying key components present at the pre-and post-synaptic levels, and understanding the mechanisms that underpin NMJ assembly. These organisms provide experimental advantages, including genetic and molecular traceability, amenability to functional studies, and increased experimental throughput compared with traditional mammalian models.

Although the NMJ of Drosophila has provided insights into the development, function and plasticity of synapses, the molecular composition and physiology of this NMJ is most similar to mammalian glutamatergic excitatory synapses and no orthologues of key molecules involved in the mammalian NMJ have been identified in Drosophila. In contrast, the use of zebrafish NMJ has continuously increased in popularity over recent decades, mainly because of the transparency of the embryonic and larval stages. This renders zebrafish particularly amenable to optical time-lapse studies in vivo. In combination with largescale genetic screens, in the late 1990s Nüsslein-Vohlhard and colleagues identified genetic mutants based on morphological abnormalities and abnormal behaviors (Granato et al., 1996; Haffter et al., 1996). Subsequent identification of the genes shed light on the molecular mechanisms underpinning the motor system; many of the locomotion mutations mapped to genes coding for synaptic proteins in the NMJs that had also been identified in mammals, such as MuSK, Dok7, Lrp4, Rapsyn or specific subunits of AChR (Marangi et al., 2001; Ono et al., 2004). These examples illustrate how these approaches complement the advances obtained using human patients or mammalian animal models, although divergence in some pathways were observed.

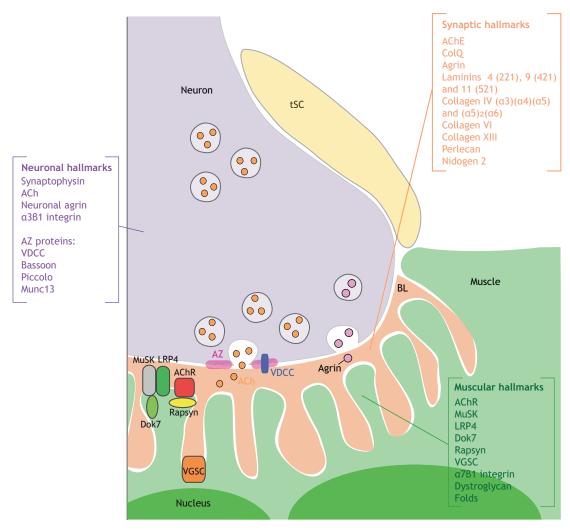


Fig. 2. Muscle, neuronal and synaptic hallmarks of the mature NMJ. The main markers that can be used to characterize the mature NMJ are shown. ACh, acetylcholine; AChE, acetylcholinesterase; AChR, acetylcholine receptor; AZ, active zone; BL, basal lamina; ColQ, collagen Q; Dok7, downstream of tyrosine kinase 7; LRP4, low-density lipoprotein receptor-related protein 4; MuSK, muscle-specific kinase; tSC, terminal Schwann cell; VDCC, voltage-dependent Ca²⁺ channel; VGSC, voltage-gated Na⁺ channel.

et al., 1998). DG is also a receptor for agrin (Gee et al., 1994), although – unlike Lrp4 – it does not activate MuSK.

The synaptic basal lamina contains many components that have important regulatory roles in myogenesis and synaptogenesis. ECM molecules help guide the process of innervation by motor neurons and are crucial to formation of the post-synaptic density as well as organizing and maintaining functional apposition of the pre- and post-synaptic elements (Maselli et al., 2012; Rogers and Nishimune, 2017). Key junctional basal lamina components include collagens [such as collagen IV $(\alpha 3)(\alpha 4)(\alpha 5)$ and $(\alpha 5)_2(\alpha 6)$ trimers, collagen XIII and ColQ), glycoproteins (such as laminin isoforms $\alpha 4\beta 2\gamma 1$, $\alpha 5\beta 2\gamma 1$ and $\alpha 2\beta 2\gamma 1$, and nidogen 2) and heparan sulfate proteoglycans (HSPGs), such as perlecan and agrin (Fox et al., 2007). The laminin β 2 chain is considered as an active zone organizer and binds to the voltage-dependent calcium channels (VDCC) located in the active zones (Nishimune et al., 2004). The interaction between laminin β2 and VDCCs leads to VDCC clustering, which is the first step toward a mature active zone. In laminin \(\beta \) knockout mice, the number of active zones and synaptic vesicles are reduced along with a reduction in neurotransmitter release. Mice lacking ColQ, collagen XIII, collagen IV $(\alpha 3)(\alpha 4)(\alpha 5)$, $(\alpha 5)_2(\alpha 6)$ trimers or collagen VI also show immature nerve terminals and/or NMJs

(Cescon et al., 2018; Sigoillot et al., 2016; Zainul et al., 2018). Networks of laminins and collagens are connected to one another by nidogen and are anchored to the sarcolemma and intracellular cytoskeleton by binding to laminin receptors: integrins and, most importantly, DG (Fox et al., 1991; Jacobson et al., 2001).

Evolution of in vitro models of NMJs

For over one century, different models and approaches have been developed to decipher the mechanisms involved in the formation, maintenance and function of NMJs. Meanwhile, increased incidence and prevalence of inherited neuromuscular diseases require the development of pertinent pathological models for deciphering the pathophysiology underlying these diseases and developing effective treatments (Bhatt, 2016). Although animal models have been crucial for the NMJ field, they have certain limitations, such as ethical issues involving animal welfare, their capacity to fully reproduce some human pathological phenotypes and in their complexity, which hinders the identification of cell-specific mechanisms, including anterograde and retrograde signals. Researchers have tried to overcome these hurdles by developing *in vitro* models to recreate the NMJs. In the following section, we focus on the contribution and the evolution of these *in vitro* models of the NMJ (Fig. 3).

A Organ explant culture **B** Dissociated explant culture* Embryonic spinal Spinal cord cord explant Axon growth Phrenic nervé - Tissue architecture Tissue architecture preserved preserved Observation of NMJs Observation of newly Diaphragm formed NMJs formed in vivo Skeletal muscle tissue explant C Mixed cell and explant culture* D Dissociated cell culture* Dissociation Tissue architecture Tissue architecture disrupted disrupted Observation of newly Dissociation Observation of newly Dissociation formed MNs, myotubes formed axons, myotubes and NMJs and NMJs

Fig. 3. Different technical approaches to studying the NMJ in vitro. (A) Organ explant culture. (B) Dissociated explant culture. (C) Mixed cell and explant culture. (D) Dissociated cell culture. (C,D) Instead of explants or dissociated cells, immortalized cell lines or motoneurons (MNs)/muscle cells generated from induced pluripotent stem cells can be used. *Chimeric model possible (different species of origin for muscle cells and MNs).

The first models of co-cultures: organ explant cultures

Historically, the first neuromuscular junction obtained in vitro was reported by R. G. Harrison in 1959 (Harrison, 1959). By cultivating medullary cord and axial mesoderm explants from clotted frog lymph between two glass slides, Harrison observed striated and contracting muscle fibers. As the frog muscle cells do not contract spontaneously, he concluded that the contractions were nerve induced. As the embryos are easy to manipulate, collect and dissect, amphibians have been extensively used as animal models for generating in vitro models of neuromuscular junctions (Anderson et al., 1977; Kidokoro et al., 1980; Peng and Nakajima, 1978; Swenarchuk et al., 1990). Among amphibians, two species stand out - Xenopus laevis and Ambystoma mexicanum - which belong to the anuran and the urodele orders, respectively. Xenopus laevis offers a number of advantages, most notably cells that are inexpensive to culture and mature rapidly (Peng et al., 1998; Swenarchuk et al., 1990). Within 3 days of co-culture, myotomal muscle cells are seen to be contracting due to innervation by neurons (Anderson et al., 1977). Cells from Ambystoma mexicanum, like Xenopus cells, can be cultivated at room temperature and in a basic cell culture medium complemented with 1% horse serum or fetal bovine serum (Anderson et al., 1977). Ambystoma cells differ from Xenopus cells in terms of the time of maturation. Ambystoma cells require a longer period to generate mature myofibers and establish functional synapses (Swenarchuk et al., 1990).

In parallel, a number of other species from chick to rodents have been used to build models for NMJs. These models were justified in part by the lack of tools to study cell biology in frog. Burgen and colleagues used rat phrenic nerve-diaphragm explants to

investigate the action of the botulinum toxin (Burgen et al., 1949) (Fig. 3A). In these first studies, models were formed by removing a section of embryonic spinal cord with the phrenic nerve and muscle still attached. These explants were maintained as 'organ cultures'.

Dissociated explants cultures and dissociated cells cultures

The next breakthrough was the development of physically separated explants at difference with the previous system (Fig. 3B). This strategy allowed the observation of newly formed NMJs (James and Tresman, 1968). In these dissociated explants cultures, most of the NMJ models aimed to visualize and understand the process of neuromuscular synapse formation and also to decipher the mode of action of pharmacological drugs (Evans, 1969). At the same time, the first serial electron microscope studies allowed detailed observations of the ultrastructure of mouse neuromuscular synapses in culture (Bornstein et al., 1968). Crain and colleagues showed that explant cultures of rat spinal cord and skeletal muscle tissues from rat and human formed functional NMJs (Crain et al., 1970).

The development of dissociated neuronal cell culture, where isolated neurons were grown on a single layer along with the availability of inverted microscopes, facilitated electrophysiological recordings *in vitro* (Scott et al., 1969). Using these tools, Fischbach and colleagues demonstrated that co-cultures of cells dissociated from chick skeletal muscle with those from chick spinal cord produced functional synaptic contacts (Fischbach, 1970) (Fig. 3C). These interspecies models, also called chimeric models, confirmed that NMJ formation and function is conserved among a variety of species.

Chimeric cell cultures

Chicks, frogs and rodents are the most common species used as sources of tissues for studying NMJ formation. Chicks offer easy access and manipulation of the embryo (Frank and Fischbach, 1979; Johnson et al., 1981), and have been used in chimeric models with rat and mouse cells. However, as mammals, the rodents *Mus musculus* (mouse) and *Rattus norvegicus* (rat) provide greater insight into human development or pathology than amphibians or birds (Ionescu et al., 2016). Mouse myoblasts can differentiate into fused and striated myofibers 2-4 days after plating (Ionescu et al., 2016). Functional synapses form 4 days later when they are cultured with murine neuronal explants (i.e. 2 days for contact between axons and myotubes followed by 2 days for synapse maturation) (Vilmont et al., 2016) (Fig. 3D). However, mouse co-cultures are rather inefficient systems, with few synapses forming, in contrast to frog co-cultures (Sanes and Lichtman, 1999).

Chimeric cell cultures models have provided answers to the origin – muscle or nerve – of a number of molecules expressed at the NMJ. For example, species-specific antibodies, one targeting specifically the HSPG from anuran species and the second targeting the HSPG from urodela species, have identified that HSPGs originate from the muscle (Swenarchuk et al., 1990). In the same vein, the development of transgenic animals brought new tools in the field of *in vitro* NMJ modeling. For example, *Hb9::GFP* transgenic mice express GFP in differentiated motoneurons (Wichterle et al., 2002). The use of *Hb9::GFP* mouse cells allow for easy visualization and identification of motoneurons in culture (Ionescu et al., 2016; Umbach et al., 2012).

Limitations and next challenges for dissociated cell models

Embryonic spinal cord explants provide motoneurons in situ, within their 3D multicellular environment (Ionescu et al., 2016). Alternatively, dissociated cells from embryonic spinal cord or myotome allow one to enrich specifically for motoneurons or muscle cells (Fischbach, 1972; Nakajima et al., 1980). Clonal and immortalized cell lines are often used to provide a large, convenient and homogenous population of cells ready to culture. However, these cells may harbor abnormal phenotypes that are not completely representative of healthy cells in vivo, either because of the immortalization process they have been through or the fact that they are cancer cells. For example, G-8 clonal myogenic cells have a modal chromosome number of 75 instead of 40 (Christian et al., 1977). Nevertheless, muscle cell lines are used largely to study the mechanism of post-synaptic differentiation and more specifically, the mechanisms of AChRs aggregation. For example, C2C12 cells are a subclone originally isolated by Blau (Blau et al., 1983) from a murine cell line generated in 1977 by Yaffe and Saxel (Yaffe and Saxel, 1977). C2C12 cells are useful to study early muscle differentiation, including myoblast fusion as well as AChR aggregation. However, C2C12 cells do not fully differentiate and do not form aggregates of acetylcholinesterase (AChE). A more recently established murine muscle cell line called MLCL forms AChE clusters in addition to the AChR clusters (Cartaud et al., 2004). Finally, collaborations between biologists and physicists lead to the emergence of new culture devices such as microfluidic devices with the objectives of better modeling NMJ pathologies, as well as providing a more suitable system for some mechanistic studies (Machado et al., 2019; Uzel et al., 2016; Zahavi et al., 2015). These developments are discussed below.

Criteria for characterizing in vitro NMJs

The establishment of reliable protocols for co-cultures led many groups to consider developing humanized *in vitro* models of NMJs

to study human neuromuscular diseases. Whichever *in vitro* model is used, a number of criteria have to be met to ascertain that an NMJ model is representative of those *in vivo* in terms of structure and function, and therefore a valid model. The criteria list that has been used is described in the next few paragraphs and is summarized in Tables 1 and 2.

NMJ morphology

In the first instance, different presynaptic, synaptic and post-synaptic hallmarks can be used as physiological and morphological clues indicative of normal NMJs. Some of them can be detected by immunolabeling of key synaptic proteins (Das et al., 2007; Dorchies et al., 2001; Umbach et al., 2012; Vilmont et al., 2016) (Fig. 2). For skeletal muscle, the first indication of complete differentiation is the morphology of the myofibers. NMJ morphology is dependent on the species as well as the stage of NMJ differentiation and the innervated muscle. Gillingwater and colleagues developed a standardized 'NMJ morph' tool to study NMJ morphology from tissue samples, highlighting these specific parameters in function of the species (Jones et al., 2017; Jones et al., 2016). The best approach to ascertain the presence of a synapse is through electron microscopy (EM). EM allows the visualization of a contact between axon terminals and muscle cells, the presence of synaptic vesicles in the presynaptic element and post-synaptic densities apposed to presynaptic elements (reviewed by Slater, 2017). However, it remains to be shown that this synapse is functional; i.e. muscle contractions induced by nerve activity and the presence of spontaneous/evoked transmission.

Muscle contractions

Striations and spontaneous contractions are also characteristics acquired during differentiation of myoblasts into myotubes in mammals. Mouse, rat and human muscle cells can contract in aneural mono-culture, unlike amphibians (Guo et al., 2014). These contractions are helpful in assessing the differentiation of the cells. Generally, two types of muscle cell contractions can be observed in co-culture models: small twitches independent of innervation (which are short in length and time) and contractions induced by innervation, which are longer and stronger (Ionescu et al., 2016). Innervation-dependent contractions are also characterized by their synchrony from the innervation of several myotubes by axonal branches of the same motoneuron (Vilmont et al., 2016). These two types of twitching movements can easily be distinguished by adding the drugs tetrodotoxin (which inhibits voltage-gated sodium channels) or d-tubocurarine (an antagonist of AChR that prevents ACh binding to its receptor) to the culture medium (Vilmont et al., 2016) (Table S1). The innervation-independent twitches are blocked only by tetrodotoxin, whereas the contractions induced by innervation are abolished by both drugs.

Functional analysis

Electrophysiological demonstration of spontaneous and evoked neurotransmission is the most detailed way to characterize the functionality of a NMJ model. mEPPS can be recorded by a microelectrode in the motoneuron and another in the myotube close to the point of contact. This criterion alone is an unequivocal evidence of a synapse, although it may not be fully mature. A functional neuron can be identified upon electrical stimulation of the motoneuron and evocation of EPPs (Nelson et al., 1993). Alternatively, EPPs can be evoked and synapses revealed by applying glutamate from a microelectrode directly onto glutamatergic motoneurons (Puro et al., 1977).

To assess whether the EPPs or contractions are dependent on neurotransmission and AChRs, different toxins or drugs can be

Fable 1. Non-human *in vitro* models of NMJ – physiological criteria

Reference	Species (muscle cells/ motoneurons/Schwann cells)	Evoked EPP/ contraction: electrical stimulus on presynaptic	Evoked contraction: glutamate iontophoresis	EPP/ mEPP	Muscle fiber contraction	EPP/contraction stopped with d-TC677.69pt or α-BTX (AChR inhibition)	EPP/contraction stop with d-TTX (VGSC inhibition)	EPP/contraction stop with agatoxin or ω -conotoxin (VDCC inhibition)	Calcium
Anderson et al. (1977)	Xenopus/Xenopus/A	Yes	ND	ND/ND	Yes	Yes	QN QN	ND	ND
Puro et al. (1977)	Rat/chick/A	ND	Yes	Yes/Yes	Yes	Yes	QN	ND	QV
Peng and Nakajima (1978)	Xenopus/Xenopus/A	ND	ND	Yes/ND	Q	ND	QN	ND	QN
Nakajima et al. (1980)	Rat/rat/A	ND	ND	ND/Yes	Q	Q.	QN	ND	QN
Kidokoro et al. (1980)	Xenopus/Xenopus /A	ND	ND	ND/Yes	Yes	Yes	QN	ND	QN
Swenarchuk et al. (1990)	Xenopus/Ambystoma/A	ND	ND	ND/Yes	Occasional	Yes	QN	ND	QN.
	or Ambystoma/rat/A				or Yes				
Nelson et al. (1993)	Mouse/mouse/A	Yes	ND	Yes/ND	Yes	Yes	Yes	ND	QN
Larkin et al. (2006)	Rat/rat/A	Yes	ND	ND/ND	Yes	Q	QN	ND	QN
Das et al. (2007)	Rat/rat/A	ND	ND	Yes/ND	Yes	Not for all myotubes	QN	ND	QN.
Umbach et al. (2012)	Mouse/mouse/A	Yes	ND	Yes/Yes	Yes	Yes	QN	ND	QN
Chipman et al. (2014)	Chick/mouse/A	ND	ND	Yes/Yes	Yes	Yes	Yes	Diminution	QN
Zahavi et al. (2015)	Mouse/mouse/A	ND	ND	ND/ND	Yes	Q	Yes	ND	Synchronized
									Ca ^{2*} signals
lonescu et al. (2016)	Mouse/mouse/A	QN	ND	ND/ND	Yes	ND	Yes	ND	Synchronized
									Ca ² * signals
Vilmont et al. (2016)	Mouse/rat/rat	ND	ND	Yes/Yes	Yes	Yes	Yes	ND	Q
Uzel et al. (2016)	Mouse/mouse (ES cells)/A	Yes (optogenetic)	Yes	ND/ND	Yes	Yes	ND	ND	Q
Machado et al. (2019)	Chick/mouse (ES cells)/A	Yes (optogenetic)	ND	ND/ND	Yes	Yes	Yes	Q	Q.
A, absent; α -BTX, α -bungard	A absent; α-BTX, α-bungarotoxin; AChR, acetylcholine receptor; d-TC, d-tubocurarine; EPP, end-plate potential; ND, no data; TTX, tetrodotoxin; VDCC, voltage-dependent Ca ²⁺ channel; VGSC, voltage-gated Na ⁺ channel	tor; d-TC, d-tubocurarine; E	EPP, end-plate pc	otential; ND,	no data; TTX, tei	rodotoxin; VDCC, voltage-dep	oendent Ca ²⁺ channe	el; VGSC, voltage-gate	d Na ⁺ channel;

used. As mentioned previously, d-tubocurarine is a toxin blocking the AChR functionally, and thus blocking twitching, EPPs and mEPPs mediated by nicotinic AChR (Chipman et al., 2014; Kidokoro et al., 1980). A common test is to record muscle cell twitching, spontaneous EPPs or evoked EPPs, and then add dtubocurarine in the medium. This addition of d-tubocurarine eliminates twitching and EPPs in the muscle cells, but not the motorneuron action potential (Umbach et al., 2012). This inhibition is reversible and when the d-tubocurarine is removed from the medium, the muscle fibers should re-start twitching or expressing mEPPs and EPPs almost immediately (Anderson et al., 1977; Das et al., 2007; Dorchies et al., 2001; Puro et al., 1977; Umbach et al., 2012; Vilmont et al., 2016).

The first humanized in vitro models of the NMJ

As the prevalence of neurodegenerative and muscular disorders is rising, and available treatments for these diseases are scarce, it is a matter of urgency to investigate new treatment strategies. Understanding the pathophysiology underlying diseases, such as amyotrophy lateral sclerosis (ALS) or congenital myasthenic syndrome (CMS), is a crucial step for early diagnosis and effective treatment. Therefore, the remaining challenge was to generate a human in vitro NMJ adapted to mimic in vivo biology with the aim to decipher pathological mechanisms and perform functional analysis.

In 1970, Peterson and Crain demonstrated that neuromuscular connections could be formed between explants of embryonic spinal cord from rodents and skeletal muscle tissue from adult humans (Crain et al., 1970). Formed with human myofibers, these cultures were maintained up to 7 weeks and electrophysiological analysis indicated that synaptic connections between rodent neurons and human muscle were functional and could be blocked with dtubocurarine. They also demonstrated an accumulation of acetylcholinesterase (AChE) at sites of synaptic contact (Crain et al., 1970). Extending this first demonstration, Peterson and Crain observed pathological features in long-term co-cultures with human dystrophic muscle. In co-cultures maintained for more than 4 months, unusual focal myofibrillar lesions in a substantial number of innervated mature Duchenne muscular dystrophy (DMD) muscle fibers were described (Peterson et al., 1986). However, the complexity of preparing these co-cultures from human muscle biopsies has restricted their use.

An important breakthrough in the field of NMJ in vitro models came with the development of co-cultures between dissociated human primary skeletal muscle cells, derived from satellite cells, with rodent embryonic spinal cords (Table 3). Kobayashi and Askanas were the first to demonstrate that monolayer cultures of adult human muscle cells can be innervated by embryonic rat spinal cord neurons and that the differentiation of human skeletal muscle cells, as well as their electrophysiological properties, are greatly enhanced by innervation (Askanas et al., 1987; Kobayashi et al., 1987). Importantly, dorsal-root ganglia were essential for achieving functional innervation, as demonstrated by the presence of regular contractions and wellorganized acetylcholinesterase-positive contacts (Askanas et al., 1987; Kobayashi et al., 1987). Further electrophysiological studies also demonstrated mEPPS and EPPs in these co-cultures were abolished by application of AChR antagonists, such as d-tubocurarine or αbungarotoxin (α-BTX), which also blocks the contractions of human myotubes in these chimeric co-cultures. These studies also highlight that innervation can occur without the original basal lamina (Kobayashi and Askanas, 1985). Finally, the development of these chimeric models of NMJ provided an important basis for pathological

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Table 2. Non-human in vitro models of NMJ - molecular criteria and summary of the studies

	Species (muscle cells/		Mature		Synaptic or				
	motoneurons/Schwann		AChR		perisvnaptic		Co-culture		Focus of the
Reference	cells)	Morphology	clusters	Pre-synaptic markers	markers	Post-synaptic markers	duration	Conclusion (Table 1 and 2)	study
Anderson et al. (1977)	Xenopus/Xenopus /A	ND	Yes	ND	QN	QN	1 week	Functional amphibians NMJ	Dvlp
Puro et al. (1977)	Rat/chick/A	Plurinucleated striated myotubes	Q	ΩN	Q	QV	3 to 10 days	Functional chimeric NMJ	Optical synapse formation
Peng and Nakajima (1978)	Xenopus/Xenopus/A	Mononuclear, striated myotubes	Yes	ND	Q	QV	few days	Functional amphibians NMJ	Dvlp
Nakajima et al. (1980)	Rat/rat/A	Multi-innervated myotubes Vesicles acc. and post- junctional folds	9	EM: active zones, vesicles	EM: Basal lamina	EM: primary and secondary folds	15 days	Functional mammalian NMJ	Dvlp
Kidokoro et al. (1980)	Xenopus/Xenopus/A	ND	Yes	ND	Q	QN Q	8 days	Functional amphibians NMJ	Dvlp
Swenarchuk et al. (1990)	Xenopus/Ambystoma/A or Ambystoma/rat/A	ND	Yes	ND	IF: HSPG	QN Q	Q	Functional amphibians NMJ	Dvlp
Nelson et al. (1993)	Mouse/Mouse/A	ND	9	ND	Q	ND	4 weeks	Functional murine NMJ	Synapse elimination
Larkin et al. (2006)	Rat/Rat/A	Multinucleated myotubes	Yes	IF: NF	QN	IF: MHC	16-18 days	Functional rat NMJ	3D NMJ model
Das et al. (2007)	Rat/Rat/A	Multinucleated myotubes	Yes	IF: NF and SYP	Q	IF: fetal myosin	12 days	Functional mammalian NMJ	NMJ model
Umbach et al. (2012)	Mouse/Mouse/A	Enlargment of the terminal bouton	Yes	IF: VaChT, ChT1, Snap25, syntaxin 1a, SYP and SV2	QN	QV	1 week	Functional murine NMJ and characterized presynaptic bouton	NMJ model
Chipman et al. (2014)	Chick/Mouse/A	Mono-innnervated myotubes	s Yes	IF: SV2, synaptic vesicle recycling (FM-4-64dye), SYP, NCAM and VDCC	QN	Q	28 days	Functional chimeric NMJ	NMJ model/ disease modeling
Zahavi et al. (2015)	Mouse/Mouse/A	Plurinucleated striated myotubes	Yes	IF: synapsin, synaptic vesicle recycling (FM-4-64 dye), Hb9 and NF-H	QN	Q	A few days	Functional murine NMJ	Dvip
lonescu et al. (2016)	Mouse/Mouse/A	Striated myotubes	Yes	IF: SYN1, SYP, Hb9 and NF	Q	QN.	3 to 4 weeks	Functional murine NMJ	NMJ model
Vilmont et al. (2016)	Mouse/rat/rat	Pretzel-like AChRs morphology, peripheral nuclei	Yes	IF: TUBB3, basoon, VaChT, ChAT, NF and SYN1	IF: GFAP (Schwann cells)	IF: AChRs, DRPR, VGSC, MuSK, rapsyn, Syne1, mFISH and £AChR	14 days	Functional rodent NMJ	Dvip
Uzel et al. (2016)	Uzel et al. (2016) Mouse/mouse (ES cells)/A	Striated myotubes	Yes	MN Hb9-GFP+	Q	IF: α -actinin	9 days	Functional murine NMJ	NMJ model
Machado et al. (2019)	Chick/mouse (ES cells)/A	Striated myotubes	Yes	IF: TUBB3, SV2 and YFP (expressed by MNs)	QN	IF: titin	9 days	Functional chimeric NMJ	Disease modeling

A, absent; acc, accumulation; DVIp, development; IF, immunofluorescence labeling; ChAT, choline acetyl transferase; ChT1, choline transporter 1; DHPR, dyhydropyridine receptor; EM, electron microscopy; Hb9, homeobox protein 9; HSPG, heparan sulfate proteoglycans; MHC, myosin heavy chain; MN, motoneuron; NCAM, neural cell-adhesion molecule; ND, no data; NF, neurofilament; RyR, ryanodine receptor; SV2, synaptic vesicle 2; SYN1, synaptophysin; TUBB3, pIII-tubulin; VaChT, vesicular acetylcholine transporter.

studies described for myotonic dystrophy (Kobayashi et al., 1990), DMD (Park-Matsumoto et al., 1991), McArdle's myopathy (Martinuzzi et al., 1993), spinal muscular atrophy (SMA), ALS and facioscapulohumeral muscular dystrophy (FSHD) (Braun et al., 1995). Several of these studies revealed some pathological phenotypes. Thus, human skeletal muscle from children affected by a severe form of SMA began to display degeneration only when cocultured with embryonic rat spinal cords (Braun et al., 1995). Despite these important advances, the only human tissue that could be used in these co-cultures was skeletal muscle cells. Recent advances in stem cell biology have fueled the prospect of generating the main components of the NMJ, offering new perspectives for pathobiological studies and therapeutic development neuromuscular diseases. Thus, the availability of human pluripotent stem cells (hPSCs), either of embryonic stem cell (ESC) or induced pluripotent stem cell (iPSC) origin, has made the generation of human muscle and nerve cells possible. The first protocols developed from hPSCs concerned the specification of spinal motoneurons. Based on developmental insights generated from animal studies and those gained on mouse ESCs (Wichterle et al., 2002), much progress has been made on the capacity to convert hPSCs into spinal motoneurons. Li and colleagues were among the first to attempt co-cultures between hPSC-derived spinal motoneurons and murine skeletal muscle cells C2C12 (Li et al., 2005) (Table 4). After 3 weeks of culture, contractions were observed only in the presence of hESC-derived motoneurons and immunostaining analysis revealed the presence of

clusters of AChR juxtaposed to neurite contacts (Li et al., 2005). Since then, this approach has been extended to disease-specific hiPSC-derived motoneurons such as SMA (Yoshida et al., 2015). In this context, the use of SMA-hiPSC derived motoneurons led to an impaired AChR clustering on C2C12 that precedes motoneuron loss (Yoshida et al., 2015). However, it is important to emphasize the relatively low rate of differentiation of hPSCs into hPSC-derived motoneurons observed in these studies, with only 25-30% of motoneurons generated on average.

To circumvent this limitation, motoneurons enriched by FACS or directly induced from fibroblasts (hi-motoneurons), without passing through an induced pluripotent stem cell stage, have been used (Du et al., 2015; Liu et al., 2016; Singh Roy et al., 2005). When cocultured with primary rodent skeletal muscle cells, both sources of motoneurons led to the colocalization of synaptic markers such as synapsin with AChR clusters. Electrophysiological analyses also revealed spontaneous end-plate currents detected in murine myotubes when co-cultured with hi-motoneurons generated from human fibroblasts by overexpression of four transcription factors (SOX11, NGN2, ISL1 and LHX3). This electrophysiological activity can be blocked by using either d-tubocurarine, tetrodotoxin or a specific inhibitor of voltage-dependent sodium channels in muscle: ωconnotoxin. When applied to a pathological condition such as ALS, profound dysfunction in the ability of patient-specific himotoneurons to control functional muscle maturation is observed (Liu et al., 2016). Although promising, these first descriptions were

Table 3. In vitro models of NMJ - human source of skeletal muscle cells

Reference	Source of MN	Co-culture duration	Pre-synaptic markers	Synaptic markers	Post-synaptic markers	Muscle contraction	EPP/ contractions blocked with d-TC or α -BTX	Disease modeling
Crain et al. (1970)	Fetal mouse spinal cord	Up to 7 weeks	ND	HC: AChE	ND	Yes	Yes	ND
Witkowski and Dubowitz (1975)	Embryonic mouse spinal cord	Up to 11 weeks	ND	ND	ND	Yes	Yes	DMD/SMA
Kobayashi and Askanas (1985)	Fetal rat spinal cord	56 days	ND	HC: AChE	IS: ¹²⁵ I-α-BTX	ND	ND	ND
Peterson et al. (1986)	Embryonic mouse spinal cord	Up to 6 months	EM: active zones and vesicles	HC: AChE	EM: primary and secondary folds	Yes	Yes	DMD Myofilament breakdown
Kobayashi et al. (1987)	Fetal rat spinal cord	Up to 4 weeks	ND	HC: AChE	IS: ¹²⁵ I-α-BTX	Yes	ND	ND
Askanas et al. (1987)	Fetal rat spinal cord	Up to 8 weeks	EM: active zones and vesicles	EM: basal lamina	EM: primary and secondary folds	Yes	ND	ND
Kobayashi et al. (1990)	Fetal rat spinal cord	Up to 7 weeks	ND	HC: AChE	IS: ¹²⁵ I-α-BTX	Yes	Yes	MD Innervation less mature
Park-Matsumoto et al. (1991)	Fetal rat spinal cord	3 weeks	ND	ND	IF: dystrophin	Yes	ND	DMD Absence of dystrophin
Michikawa et al. (1991)	Rat spinal cord	2 weeks	ND	HC: AChE	IF: α-BTX	Yes	Yes	ND
Braun et al. (1995)	Embryonic rat spinal cord	1 month	ND	ND	EM: primary and secondary folds	ND	ND	SMA (type I, II and III) Degenerating co-cultures
Dorchies et al. (2001)	Rat spinal cord	Up to 1 year	ND	ND	EM: primary and secondary folds IF: desmin, troponin- T, tropomyosin, MHC and ¹²⁵ l-α-BTX	Yes	ND	X-linked MTM No phenotype

 α -BTX, α -bungarotoxin; AChE, acetylcholinesterase; d-TC, d-tubocurarine; DMD, Duchenne muscular dystrophy; EM, electron microscopy; HC, histochemistry; IF, immunofluorescence labeling; IS, iodine staining; MD, myotonic dystrophy; MHC, myosin heavy chain; ND, no data; SMA, spinal muscular atrophy; X-linked MTM, X-linked myotubular myopathy.

Table 4. In vitro models of NMJ - human source of motoneurons

Reference	Source of skeletal muscle cells	Co-culture duration	Pre-synaptic markers	Post- synaptic markers	Muscle contraction	EPP/contractions blocked with d-TC/TTX/ω-conotoxin	Disease modeling
Singh Roy et al. (2005)	Rat myoblasts	1 week	IF: SV2, TUBB3 and MAP2	IF: AChRs and desmin	ND	ND	ND
Li et al. (2005)	C2C12	Up to 2 weeks	IF: SYN1 and ChAT	IF: AChRs	ND	ND	ND
Du et al. (2015)	C2C12	10 days	IF: ChAT	IF: AChRs	ND	ND	ND
Yoshida et al. (2015)	C2C12	Up to 60 days	IF: SV2 and NF	IF: AChRs and MHC	ND	ND	SMA type 1 defective AChR clustering
Guo et al. (2010)	Rat skeletal muscle	3 weeks	IF: SYP and TUBB3	IF: AChRs and MHC	Yes	Yes Electrophysiological recordings	ND
Liu et al. (2016)	Mouse myoblasts	1 week	IF: SYN1	IF: AChRs and MHC	ND	ND	ALS (FUS mutations) Decrease AChR clusters and defective AP (amplitude/ frequency)

AChR, acetylcholine receptor; ALS, amyotrophic lateral sclerosis; AP, action potential; ChAT, choline acetyl transferase; d-TC, d-tubocurarine; EM, electron microscopy; IF; immunofluorescence labeling; MAP2, microtubule-associated protein 2; MHC, myosin heavy chain; ND, no data; NF, neurofilament; SMA, spinal muscular atrophy; SV2, synaptic vesicle type 2; SYN1, synapsin 1; TTX, tetrodotoxin; TUBB3, β-III tubulin.

all based on chimeric models and to reach the 'holy grail' of a 'real' human NMJ *in vitro* model, both co-culture components should be of human origin.

Towards entirely humanized NMJ in vitro models

Over the past decade, the literature has documented fully humanized in vitro models of NMJs based on motoneurons mainly derived from hPSCs cultured with human primary skeletal muscle cells (Afshar Bakooshli et al., 2019; Marteyn et al., 2011; Santhanam et al., 2018; Shimojo et al., 2015; Steinbeck et al., 2016; Vila et al., 2019). Although all these studies diverged in their protocols, they all described the presence of AChR clusters in close proximity to neuritic processes, as well as muscle twitching after 2 or 3 weeks of co-culture. This indicates to some extent the functionality of human primary skeletal muscle cells when co-cultured with human motoneurons (Table 5). To go further ahead in the functional characterization of this system, Steinbeck and colleagues were the first to combine hPSCbased in vitro models of NMJs with optogenetics (Steinbeck et al., 2016). The authors used purified hESC-derived motoneurons stably expressing channelrhodopsin 2 (ChR2) coupled to the GFP under the human synapsin promoter. After 6-8 weeks of co-culture with human primary skeletal muscle, muscle twitches were observed and quantified after light stimulation. The addition of vecuronium, an antagonist of the nicotinic AChR, completely blocked the lightinduced muscle twitches. Calcium imaging and electrophysiological analyses also confirmed these results. As purification of hPSC-derived motoneurons could lead to the selection of physiologically irrelevant motoneurons, these datasets were reproduced using motoneurons generated with a recent protocol that allows the efficient conversion of hPSC into a homogenous population of spinal motoneurons without using a purification step (Maury et al., 2015). Interestingly, the authors also demonstrated that their 'all human' NMJ cultures could be used in the pathological context of myasthenia gravis (Box 1) (Steinbeck et al., 2016).

Despite these developments, several challenges remain. The first limitation is the source of the human skeletal muscle cells. All the models described are based on the use of human primary skeletal muscle cells, which exhibit substantial donor variation in terms of differentiation (Nikolic et al., 2017). The development of an 'all human' NMJ model in which both neuron and muscle components are

derived from the same individual is of great interest for modeling disease of genetic etiology. With the recent development of protocols allowing the conversion of hPSCs into skeletal muscle cells (reviewed by Pourquie et al., 2018), it is possible to produce co-cultures between motoneurons and skeletal muscle cells derived from hiPSCs from the same donor (Demestre et al., 2015; Maffioletti et al., 2018; Mazaleyrat et al., 2020; Osaki et al., 2018; Picchiarelli et al., 2019; Puttonen et al., 2015). Although the maturation level of these co-cultures generated from 'embryonic-like' pluripotent stem cells is still questionable, the detection of AChR clustering, contractions and calcium imaging suggest an acceptable level of functionality of these systems.

Another limitation stems from the fact that most of these studies have used a 2D cell culture system that does not mimic the native tissue structure. Consequently, different attempts have been made recently to evaluate the possibility of developing an 'all human' NMJ in vitro model by taking advantage of the advent of 3D cultures and microfluidic lab-on-a-chip technologies (Afshar Bakooshli et al., 2019; Maffioletti et al., 2018; Osaki et al., 2018; Santhanam et al., 2018; Vila et al., 2019). Microfluidics allows precise control over the cell microenvironment, providing an interesting platform for studying the NMJ. Motoneurons and muscle can be in two different chambers connected by axons, which enables spatiotemporal control over the cells. Analysis of the localized actions at the soma, along the axon and at the NMJ can be performed. Santhanam and colleagues were among the first to apply microfluidics to human motoneurons and muscle cells. Two weeks of co-culture were sufficient to observe muscle contractions after electrical stimulation of the motoneurons that can be blocked by treatment with d-tubocurarine (Table S1). However, the percentage of contracting myotubes is lower than the number of NMJs determined immunocytochemically, suggesting that some of the latter may not be mature enough to be functional (Santhanam et al., 2018).

Another approach relies on the use of 3D co-cultures with muscle cells in hydrogels, such as the commercially available basement membrane extract Matrigel followed by differentiating the motoneurons on the surface. 3D cultures tend to be functionally more relevant with respect to contractile muscle. Collagen I is a major component of the native ECM and harbors adequate properties needed for contraction and force generation. However, Collagen I has less stability, degrades faster and appears to be a less effective solution for

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Table 5. In vitro models of NMJ - human source of skeletal muscle cells and motoneurons

	All from		On this co		Dro cyto ordio			Missis	anditocatago/GGT	Saiscai aniolo	
Reference	hPSCs	Cell origin	duration	2D/3D	markers	Synaptic markers	Post-synaptic markers	contraction	blocked with drugs		Disease modeling
Guo et al. (2011)		Human primary myoblasts and hESC-derived MNs	2-3 weeks	2D	IF: SYP and TUBB3	QN	IF: AChRs and MHC	Yes	Yes	QN	ND
Marteyn et al. (2011)		Human primary myoblasts and hESC-derived MNs	2 weeks	2D	IF: TUBB3	Q	IF: AChRs	Q	Q	QN QN	DM1 (defect in neuritic arborization)
Puttonen et al. (2015)	Yes	hESC mixed differentiation	Up to 6 weeks 2D	2D	IF: SV2, SYP	Q	IF: AChRs, MHC and	Yes	Yes	Under stimulation with	ND
Steinbeck et al. (2016)		Human primary myoblasts and hESC-derived MNs	6-8 weeks	2D	IF: SYP	Q	IF: AChRs, desmin and MHC	۶	Yes, after optogenetic	ChR2	MG (decreased muscle twitches)
Osaki et al. (2018)	Yes	hiPSC-derived muscle cells 2 weeks and hiPSC-derived MNs	2 weeks	3D	IF: ChAT, SMI-32, TUBB3	ND	IF: AChRs, α-actinin	Sumulation Yes	Stimulation	ChR2 (CAG promoter, AAV transduction)	ALS (degenerescence, fewer contractions)
Maffioletti et al. (2018)	Yes	hiPSC-derived muscle cells, hiPSC-derived MNs and hiPSC-derived	2 weeks	3D	IF: SMI-32	Ŋ	IF: AChRs, MHC and Titin	ND	Ŋ	ON CONTRACTOR	Laminopathies Defect in nuclear elongation
Santhanam et al. (2018)		Human primary myoblasts and hiPS-derived MNs	2 weeks	2D microfluidic	IF: NF and SYP	QN	IF: AChRs and MHC	Yes	Yes	QN	ND
Afshar Bakooshli et al. (2019)		Human primary myoblasts and hESC-derived MNs	2 weeks	2D versus 3D	IF: SYP and SMI-	Laminin β2	IF: AChRs, α-actinin and rapsvn	Yes	Q	GCaMP6/ChR2 (EiFa	MG (decreased area of muscle response)
Vila et al. (2019)		Human primary myoblasts and ChR2-expressing hiPSC-derived MNs	3 weeks	3D microfluidic	Q Q	Q	IF: œactinin	Yes	Yes	ChR2 (EiFa promoter)	MG (difference in contractile properties)
Lin et al. (2019)	Yes	hiPSC-derived muscle cells Up to 100 and hiPSC-derived MNs days	Up to 100 days	2D	IF: TUBB3, NF, SV2 and ChAT	QN	IF: AChRs and MHC	No contraction without stimulation	QN	ChR2 (synapsin promoter)	SMA (smaller area of NMJ)
Picchiarelli et al. (2019)	Yes	hiPSC-derived muscle cells and hiPSC-derived MNs	3 weeks	2D	Q	Q	IF: AChRs and α- actinin	Q	Q	QN QN	ALS (defect in AChR clustering)
Bellmann et al. (2019)	Yes	hiPSC-derived muscle cells and hiPSC-derived MNs	3 weeks	3D microfluidic	IF: TUBB3, ChAT and synapsin EM: vesicles and primary folds	ND	IF: AChRs and α- actinin	No contraction without stimulation	ND	ChR2 (EiFa promoter)	QN
Faustino-Martins et al. (2020)	≺es	hiPSCs and hESCs	Up to 150 days	3D organoids	IF: ChAT, SMI-32, TUBB3, SYP, GFAP, S100β and MBP EM: synaptic cleft and vesicles	IF: laminin sheaths	IF: fast MHC and AChRs EM: sarcomeric units	Spontaneous Blocked by curare	Blocked by TTX	Calcium signal MEA	MG (severe reduction of NMJ)
Kim et al. (2020)		hMPCs and hNSCs	10 days	3D bioprinting	IF: TUBB3, NF and GFAP	Q	IF: MHC AChRs	Q	Q	Calcium signal	Rodent muscle defect injury model
Mazaleyrat et al. (2020)	Yes	Concomitant differentiation of hiPSC-derived muscle cells and hiPSC-derived MNs	30 days up to 7 months	2D	IF: NF and S100ß	Q	IF: desmin, Pax3/7, titin, MHC and AChRs EM: sarcomeric units and peripheral	Spontaneous	Blocked by α-BTX and TTX	Calcium signal	DMD, DM1, FSHD2 and LGMD2A (multiple alterations)

α-BTX, α-bungarotoxin; ACh, acetylcholine; AChR, acetylcholine receptors; ALS, amyotrophic lateral sclerosis; BTX, bungarotoxin; ChAT, choline acetyl transferase; ChR2, channelrhodopsin 2; d-TC, d-tubocurarine; DM1, myotonic dystrophy type 1; EM, electron microsoopy; GFAP, gial fibrillary acidic protein; IF, immunofluorescence labeling; hMPCs, human mesenchymal precursor cells; hNSCs, human neural stem cells; hPSCs, human pluripotent stem cells; MBP, myelin basic protein; MG, myasthenia gravis; MHC, myosin heavy chain; ND, no data; NF, neurofilament; SV2, synaptic vesicle type 2; SYP, synaptophysin; TTX, tetrodotoxin; TUBB3, β-III tubulin.

long-term cultures (Leikina et al., 2002). In contrast to Collagen I, Matrigel seems to be better adapted for muscle differentiation (Grefte et al., 2012). Other alternatives to collagen are fibrin, fibrinogen or thrombin (Wu et al., 2005). Such approaches have recently been applied to the development of fully humanized NMJ in vitro models in which muscle cell laden hydrogels were formed in specific molds with pillar structures at each extremity sufficient. These then created a continuous uniaxial tension and directed orientation of cells along the axis of tension (Afshar Bakooshli et al., 2019; Maffioletti et al., 2018). A comparative analysis of 2D versus 3D co-cultures reveals a faster and more efficient functional innervation in 3D versus 2D co-cultures when reviewed after 2 weeks in culture. The determination performed using calcium imaging, optogenetics and electrophysiological recordings (Afshar Bakooshli et al., 2019). A more mature expression profile of AChR isoforms was observed in 3D co-cultures with an increased expression of the adult AChR ε isoform in comparison with 2D cultures. Nonetheless, the small size of muscle fibers, as well as the properties of action potentials recorded in these co-cultures when compared with in vivo mammalian models, suggests that the 3D human NMJs co-cultures are not fully mature and that additional cues might be necessary to ensure further maturation of tissues (Afshar Bakooshli et al., 2019). Increasing culture time, adding trophic factors or providing other cell types have been suggested. Recently, complex isogenic multilineage 3D co-cultures have been developed containing skeletal muscle cells, vascular endothelial cells, pericytes and motoneurons, all derived from hiPSCs (Maffioletti et al., 2018). Whereas these complex co-cultures were mainly characterized at the morphological level, these results open new perspectives for the development of humanized organoid-like platforms for studying NMJ development and disease modeling.

Two studies have also described the combination of 3D system and microfluidics (Osaki et al., 2018; Vila et al., 2019). These approaches have the advantage that motoneurons and muscle cells can be compartmentalized while providing spatial guidance cues and signal gradients using 3D hydrogels for better integration. The study uses engineered functional contractile hiPSC-derived skeletal muscle cells co-cultured with hiPSC-derived motoneurons. Interestingly, such approaches are compatible for studying and validating the therapeutic potential of compounds for neuromuscular disorders such as ALS (Osaki et al., 2018).

Finally, 3D bioprinting is an emerging field that could be relevant for the development of *in vitro* NMJ systems. A recent study described the feasibility of using bio-printed 3D hydrogel skeletons to support culture, differentiation and alignment of human primary skeletal muscle cells compatible with the functional integration of immortalized human neural progenitor cells (Kim et al., 2020). Interestingly, these bio-printed constructs with neural cells facilitate rapid innervation and mature into organized muscle tissue, restoring normal muscle function in a rodent model of muscle defect injury (Kim et al., 2020). This development can be pursued in the future for improving the maturity of NMJ systems and highlights an area of considerable interest in the tissue-engineering field.

Future perspectives

For over one century, the field of research around NMJs has grown constantly. Recent advances in stem cell biology have stimulated the development of innovative models of human NMJs. Despite initial successes in developing human motoneuron-muscle cells co-cultures, stem-cell derived NMJs remain immature in comparison with the human and chimeric NMJ models. One challenge is the limited survival of these co-cultures. The first chimeric mouse-human co-cultures could survive for more than 6 months (Peterson et al., 1986),

whereas human stem-cell based co-cultures often do not survive beyond 4 weeks. However, a recent protocol of human stem cell codifferentiation enables the co-culture to survive up to 7 months (Mazaleyrat et al., 2020). Another challenge for future studies is to find a different method for generating spinal motoneurons. Among the motoneuron lineages in vivo, seven different lower motoneuron subtypes exist, each of them innervating a different muscle group. Most of the NMJ models from iPSCs in the currently published studies used iPS cell-derived general motoneurons. Efforts are under way to characterize the subtypes obtained and to improve the motoneuron differentiation protocols in order to obtain specific motoneuron subtypes (Hester et al., 2011; Lukovic et al., 2017). Similarly, iPS cell-derived muscle fibers are so immature that they cannot be classified as muscle fiber type I or II. Future work to obtain well-characterized motoneurons and muscle fibers subtypes will be particularly important when it comes to modeling neuromuscular pathologies, as all NMJs are not affected in the same way (Nicole et al., 2014). Additionally, tSCs are still a missing partner in current in vitro models. Protocols enabling the differentiation of iPS cells into myelinated Schwann cells are available (Liu et al., 2012; Ziegler et al., 2011). To date, there are no protocols for generating tSCs. These cells, together with complex and specialized basal laminas, will likely be required to reach a fully mature in vitro human NMJ model. Future research on 3D in vitro human NMJ models will be important in this effort. Recently, Faustino Martins and colleagues developed an elegant strategy to derive neuromuscular organoids from hiPSCs. Over a period of 50 to 100 days, these organoids reached a size of 5-6 mm in diameter and displayed an elongated morphology with neural tissue on one side and muscle cells on the other. In this system, electron microscopy confirmed the presence of presynaptic nerve terminals containing synaptic vesicles and active zones: tSCs and post-synaptic muscle fibers with organized sarcomeres and junctional folds. Importantly, these NMJs were functional, as demonstrated by the detection of muscle contraction after 50 days of culture. The functionality could be inhibited with d-tubocurarine, as well as spontaneous calcium oscillations and electrical activities (Faustino Martins et al., 2020). When employed with patient-specific stem cells together with CRISPR/Cas9 genome editing to restore or create mutations identified in patients, these new models will provide more reliable models of human NMJs in vitro and of neuromuscular diseases. Finally, to address the needs of new therapeutic approaches for neuromuscular diseases, standardized and miniaturized human in vitro NMJ models are a long-awaited drug-screening tool.

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Competing interests

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