Guiding principles for drug discovery and development in amyotrophic lateral sclerosis

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Abstract

The level of research activity undertaken globally into the causes of amyotrophic lateral sclerosis has dramatically expanded in recent years, resulting in the identification of multiple genetic risk factors, cellular pathways, and potential drug targets. Despite these advances, ALS remains an inexorably progressive and fatal disease with limited therapeutic options. This expert led review is the product of a series of focussed workshops with participants from basic and translational ALS research groups, industry professionals and research funding agencies. The outputs of these workshops were then introduced to international ALS stakeholders to comment via a Delphi survey and through an additional online workshop engaging over 60 experts globally, to further refine and shape the guidance. We discuss the factors governing the barriers to translation of advances in basic science into an effective therapy for ALS, and suggest guiding principles for de-risking clinical trials of novel therapeutics by an enhanced evidenced-based approach to preclinical studies.



Authorship

This white paper was developed by an expert working group which met throughout 2022. The group was convened by the MND Association and My Name'5 Doddie Foundation and coordinated by Medicines Discovery Catapult.

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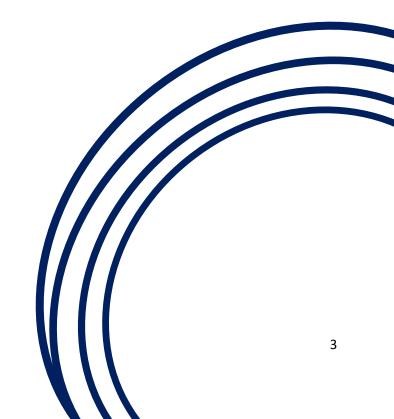
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Introduction

Amyotrophic lateral sclerosis (ALS) is a neurodegenerative disease characterized by progressive paralysis, ultimately resulting in respiratory failure and death, on average 2-3 years from first reported loss of function, but with wide variation. ALS has a median onset of 65 years and a lifetime risk of 1/350, in European populations ⁽¹⁾⁽²⁾. A reliable diagnosis can be made using clinical criteria and supportive neurophysiological tests once the disease is well established, though early diagnosis remains a challenge ⁽³⁾. Our understanding of the complex biology of ALS, which is characterized by considerable clinical, genetic and pathological heterogeneity is still incomplete, ⁽⁴⁾⁽⁵⁾. Little is known about the biological basis for the variation in clinical features such as rate of disease progression.

Factors leading to high failure rates in ALS clinical trials include the inadequacy of preclinical models in predicting efficacy in humans, significant delay in diagnosis, and uncertainty about which pathways define the most treatable phase of the disease. Only one drug, riluzole, has been approved for ALS treatment by multiple regulatory agencies internationally.

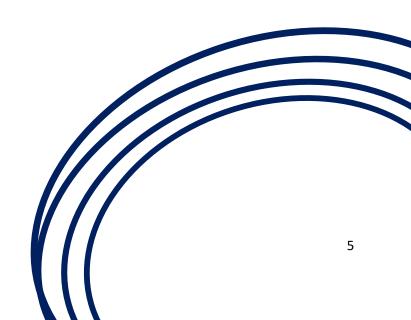
Building on existing recommendations for preclinical studies in rodents, and for the design and implementation of ALS clinical trials ⁽⁶⁾⁽⁷⁾, a cross-sector group of ALS experts in academia and industry have combined to i) agree a consensus on the key reasons for poor translational outcomes in ALS, and ii) generate recommendations for the optimization and conduct of preclinical studies to improve confidence that a drug will show efficacy in human trials. It is recognized that every therapeutic programme will be different and may employ some but not all of these principles. This guidance does not intend to stifle innovation, but to catalyse high quality preclinical drug discovery efforts and create a shared vision for all stakeholders involved in ALS drug discovery research.



Factors responsible for poor translation in ALS drug discovery and development

The persistent high failure rate in ALS clinical studies, despite decades of ALS drug discovery efforts, is due to a combination of factors outlined below.

Comp	lex Aetiology and Pathogenesis
• Procli	Incompletely understood, including relative importance of early versus late events nical models do not predict efficacy in human trials
• • • • • • • •	Lack of implementation of existing guidance ⁽⁷⁾⁽⁸⁾ Historical reliance on SOD1 animal models (non-TDP-43 pathology) Models mimic limited aspect of ALS pathology and lack a human biological context (e.g. TDP-43 regulation of cryptic exon splicing) Emphasis on survival as an endpoint, particularly in SOD1 mice ⁽⁷⁾ No credible models for 'sporadic' disease No reliable large animal models (relevant for CSF biomarkers, PK and other parameters) Lack of independent verification between laboratories or through CROs, and across different models Failure to demonstrate target engagement Lack of biomarkers correlating with efficacy in preclinical models Failure to align animal-model findings with human data (genetics, tissue, fluid samples) Lack of effective PK/PD to inform human dosing imental Medicine is underdeveloped
•	Weak rationale derived from other neurodegenerative diseases (i.e., for small molecule repurposing) Lack of data for blood-brain barrier penetration, target engagement, biomarker response, dose finding No framework for stratification according to biological readouts (genotype, pathway specific biomarkers) al studies are onerous, time-consuming, and expensive
•	Lack of robust measures of disease progression, limitations of ALSFRS-R as primary endpoint Need for better clinical and patient-reported outcome measures, and statistical powering Lack of stratification/enrichment strategy by drug mechanism Trials historically performed late in disease due to delayed diagnosis Short study length may not predict long-term benefit Prohibitive cost with current trial designs, need for digital endpoints and PPIE in trial design



Therapeutic case studies

To better understand how to develop best practice in ALS drug discovery and development, we have considered several key therapeutics that have succeeded or failed in clinical trials as case studies.

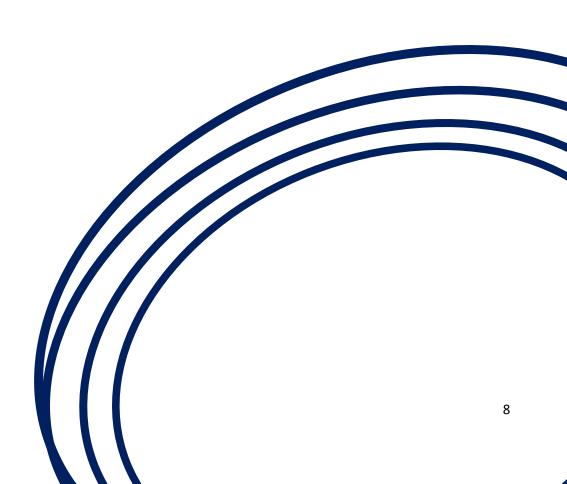
Ration	ale:
pharma glutama	
Preciin	nical studies:
No pre	clinical studies were conducted prior to the first riluzole clinical study in ALS.
Clinica	al studies:
•	The first randomized controlled trial demonstrated a modest increase in survival at one year with riluzole compared to placebo ⁽¹¹⁾ . A much larger dose-ranging study confirmed a small but statistically significant increase in survival with intermediate and high dose riluzole ⁽¹²⁾ . A Cochrane review of 4 riluzole clinical studies concluded that riluzole prolongs median survival by about two to three months in people with ALS ⁽¹³⁾ .
Implica	ations for future studies:
means using b	e has multiple pharmacological effects. A lack of understanding of the exact mechanism of action in ALS there is no biomarker and patient stratification strategy. Systematic prospective or retrospective studies blood and CSF neurofilaments as potential treatment response markers are not available. No quality-of- a is available for riluzole.



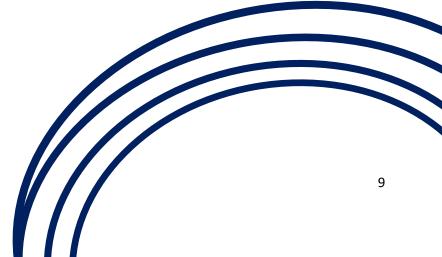
Edaravone, a free radical scaven	ger, is licensed for the treatment of acute ischaemic stroke in Japan. Excess
	inked to neuronal damage in ALS.
Preclinical studies:	
Efficacy: SOD1 G93A mice, intraperitoneally administered edaravone upon identification of clinical onset ⁽¹⁴⁾ . Efficacy: SOD1 H46R rat model	 Modest effect on behavioural function and motor neuron sparing in these models No effect on survival was observed No significant change in behavioural function or survival
⁽¹⁵⁾ . Efficacy: Wobbler mouse, Intraperitoneally administered with edaravone upon Identification of clinical onset ⁽¹⁶⁾ .	 Higher dose (10 mg/kg) of edaravone treatment significantly attenuated muscle weakness and contracture in the forelimbs, and suppressed denervation atrophy in the biceps muscle and degeneration in the cervical motor neurons compared to vehicle. The study was limited to a short duration of only 4 weeks.
Clinical studies:	
 treatment with edaravone better with treatment (17). evidenced by a smaller de evidenced by a smaller de A subsequent phase 3 studefined subset of patients life deterioration compare 	no significant delay in disease progression as measured by ALSFRS-R upor . However, one of the secondary endpoints, pinch strength, was significantly A post-hoc analysis did identify efficacy in a small subset of people with ALS ecline in ALSFRS-R score compared to placebo ⁽¹⁸⁾ . udy attempted to investigate the safety and efficacy of edaravone in this mor and did demonstrate less functional loss at 6 months and reduced quality o d to those receiving standard of care ⁽¹⁹⁾ . y aimed to evaluate the long-term safety and effectiveness of intravenous
edaravone therapy for pa 116 patients treated for a treated for a median of 11 term intravenous edaravo	tients with ALS in a real-world clinical setting. Disease progression among median of 13.9 months with edaravone did not differ from 116 patients 1.2 months with standard therapy. The study concluded that although long- one therapy for patients with ALS was feasible and mainly well tolerated, it any disease-modifying benefit ⁽²⁰⁾ .

Case study: tecfidera (dimethyl fumarate)			
Rationale:			
Tecfidera is licensed for the treatment of multiple sclerosis, where it was shown to alter the immune response toward a more anti-inflammatory profile by increasing the number of regulatory T-cells. Alterations in regulatory T lymphocytes correlate with disease progression rate and severity in ALS ⁽²¹⁾ , supporting the hypothesis that Tecfidera would reduce inflammation and potentially slow disease progression in people living with ALS.			
Preclinical studies:			
No ALS preclinical studies have been published, to our knowledge.			
Clinical studies:			
In a phase 2 study, treatment with Tecfidera failed to show a significant difference in ALSFRS-R score at week 36 and there were no significant changes in other secondary outcome measures (survival, neurophysiological index, respiratory function, urinary neurotrophin-receptor p75 and quality of life) ⁽²²⁾ .			
Implications for future studies:			
 Preclinical studies in relevant ALS models (<i>in vivo</i> and/or patient-derived cells) are recommended before progression to clinical study, to demonstrate mechanism of action, efficacy, predict dose for human studies and identify relevant biomarker or patient stratification strategy. 			

• Proof of mechanism studies in ALS patients could also be valuable to demonstrate back translation from patients to preclinical models.

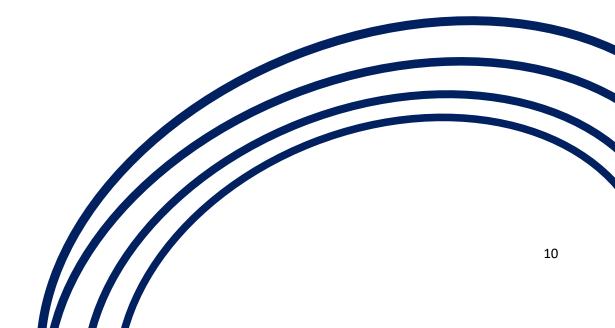


Rationale:	
Gain of function genetic variants in SOD1 are a oligonucleotides provide a method to directly ta Preclinical studies:	associated with 20% of familial ALS cases. Antisense arget gene expression
Target engagement: by ASO administration in SOD1 G93A mice and rats ⁽²³⁾ . Efficacy: SOD1 G93A mice and rats, by injection of Tofersen at 50 and 94 days, then 80 and 110 days ⁽²³⁾ .	 CNS penetration determined Reduction of SOD1 expression achieved Tofersen treated mice maintained weight, performed better on the rotarod assay and had extended survival
Markers of disease: were studied after dosing SOD1 G93A at 5 weeks of age ⁽²³⁾ .	 Compound muscle action potential (CMAP) and innervation of the tibialis anterior muscles in the hind limbs was maintained in treated mice Treated mice showed lower levels of serum phospho- neurofilament heavy chain (pNFH) levels
Reversal of SOD1-mediated neuronal dysfunction: Mice were treated at 9 weeks of age when CMAP had clearly begun to decline and pNFH had begun to rise ⁽²³⁾ .	Substantial increase of CAMP and decrease in pNFH was seen in treated mice
Proof-of-principle in non-human primates (cynomolgus monkeys) (23).	 Demonstrated successful reduction in SOD1 expression and determined PK/PD relationship
Clinical studies:	
 Tofersen administered intrathecally over Phase 3 failure: after 28 weeks of thera improvements in the primary endpoint Functional Rating Scale (ALSFRS-R) or 	apy, patients did not demonstrate statistically significant assessed by the Revised Amyotrophic Lateral Sclerosis compared to placebo. However, preliminary data suggests that wed signs of slowing disease progression, particularly in
	studies that were late on in disease progression was
 inconsistent with the state of disease p treatment effect. Possibility for more sophisticated precl pathway modulation, motor unit surviva 	studies to identify a patient stratification strategy and identify



Case study: ozanezumab

Rationale:				
identified as a potent myelin-as	Nogo-A), a high molecular-weight transmembrane protein, initially ssociated inhibitor of axonal growth, is upregulated in ALS and might promote specific antibody directed against Nogo-A.			
Target engagement: in SOD1	Co-localisation of ozanezumab with Nogo-A between muscle fibres			
G93A mice ⁽²⁷⁾	 Co-localisation of ozariezumab with Nogo-A between muscle libres and tissue via immunohistochemistry, and increased Nogo-A RNA levels 			
Efficacy: Symptomatic treatment of SOD1 G93A mice from 70 days of age with Ozanezumab ⁽²⁷⁾	 Compared with vehicle-treated SOD1G93A mice, early symptomatic treatment with anti-Nogo-A antibody from 70 days of age significantly improves muscle innervation, muscle function and phenotype and increases motor neuron survival in symptomatic mice SOD1G93A mice at 90 days of age However, the improvement in muscle function and motor neuron survival was not maintained and by 120 days most of the beneficial functional effects of anti-Nogo-A treatment was no longer observed in SOD1G93A mice 			
Clinical studies:				
 In a Phase 2 trial, Oza primary outcome was a 	tolerated in a first-in-human trial ⁽²⁸⁾ . nezumab did not show efficacy compared with placebo in patients with ALS. The a joint-rank analysis of function (ALSFRS-R) and overall survival, analysed at 48 ho received at least one dose of study drug ^{(29).}			
Implications for future studie	es:			
 Preclinical studies indicated that treating early with ozanezumab during disease progression may represent a valid therapeutic window to delay disease progression, yet patients were not stratified by stage of disease progression in clinical studies. Future studies should incorporate patient stratification approaches, including stage of disease, where appropriate. 				



Rationale:	
acid) are proposed to reduce neu and mitochondrial dysfunction. Preclinical studies: Preclinical data support a mitigat death and other disease-specific dysfunction. However, no publish	s sodium phenylbutyrate and taurursodiol (also known as tauroursodeoxycholic uronal death in ALS by simultaneously mitigating endoplasmic reticulum stress ting effect of sodium phenylbutyrate (PBA) or taurursodiol alone on neuronal features in models of neurodegenerative diseases and mitochondrial ned preclinical studies of Taurursodiol undertaken specifically in relation to reclinical evidence of Sodium Phenylbutyrate-Taurursodiol treatment in ALS
models.	
Efficacy of PBA: In G93A SOD1 mice treated 21 before symptoms occur ⁽³⁰⁾ and in combination with riluzole (del Signore et al., 2009).	 Compared to vehicle-treated SOD1 G39A mice, treatment with PBA resulted in extended survival, improved motor performance as demonstrated by the rotarod assay, improved stride length and body weight and reduced hindlimb muscle wasting In combination with riluzole, PBA showed synergistic effect in improving body weight loss, grip strength deficit and life span.
Functional assays PBA: in G93A SOD1 mice treated 21 before symptoms occur ⁽³⁰⁾ and in combination with riluzole ⁽³¹⁾	 PBA demonstrated a neuroprotective effect, as demonstrated by reduced atrophy of the lumbar spinal cord and ventral neuron loss compared to untreated mice PBA also prevented ubiquitinated-positive aggregates in spinal cord ventral horn neurons Molecular mechanism of PBA determined and PBA shown to restore dysregulated signalling in ALS mice. In combination with riluzole, PBA attenuated lumbar ventral horn neuronal loss and reactive astrogliosis by increasing Histone 4 acetylation and NF-kBp50 translocation to the nucleus in G93A mice
Clinical studies:	
 ALS ^(32,33). In a phase 2 trial, Sodiur placebo as measured by were significantly difference neurofilament H subunit In an open-label extension 	
models.	of preclinical findings from other diseases should be carried out in ALS extending study duration beyond six-months to identify long-term effects on



Guiding principles for ALS drug discovery and development

Taking into account failures of pivotal ALS clinical studies and best practice in the field, we propose a framework of guiding principles for ALS drug discovery and development programmes, with the aim of increasing confidence in a new target, drug and clinical study design to enable the best chance of success in clinical trials. Given that we do not yet have a definitive set of principles for successful ALS drug discovery and development, we suggest that a pragmatic view must be taken on a case-by-case basis to ensure that frameworks such as this do not slow progress

A key element required to enable the implementation of these guiding principles, is the provision of appropriate funding. Some of the studies outlined are costly but do not fit within traditional grant funding schemes e.g., funding for independent validation of preclinical efficacy or Experimental Medicine studies. Integration of resources and reducing barriers to accessing appropriate infrastructure are also fundamental requirements to enable innovators to implement the principles outlined below.

Disease biology and linkage	Definition of the drug target, and its relation to ALS pathophysiology	releva linka	int ti ge ti	of target in issue and o ALS in confirmed	m mod	tage of disease for nost effective target dulation hypothesised symptomatic vs pre- symptomatic
Target validation	Develop platform of evidence for target relevance, differentiation, and target safety	Target engagement and downstream pharmacodynamic biomarkers defined		Target modulation shown to result in pathway modification and rescue of disease phenotype <i>in vitro</i> e.g., iPSC or other relevant cellular model		
Preclinical studies	 PD studies - Drug shown to modulate functionally relevant pathway <i>in vitro</i> <u>and/or</u> <i>in vivo</i> (e.g., iPSC models, mouse model where target pathway is dysregulated and/or <i>in vivo</i> model of ALS). Build body of evidence by repeating a hypothesis using multiple relevant models or approaches 		Target tissue exposure and target engagement demonstrated <i>in</i> <i>vivo</i> (e.g., wild- type mouse)		PK studies in wild-type mice; determine appropriate dosing regimen and route of administration	
Experimental Medicine	Target tissue exposure and target engagement demonstrated Understanding of effect of drug on downstream pathways and disease biology			Evidence functiona response	d	Robust biomarker and patient stratification strategy established

Use of preclinical models of ALS in drug discovery and development

In contrast to the successful use of preclinical models in developing antisense oligonucleotide therapy (in ALS and SMA), there is currently no example of a preclinical model which has successfully predicted efficacy of a drug in clinical trials in sporadic ALS. Until this has been achieved, proposing one or more 'ideal' preclinical models remains aspirational. However, a number of promising preclinical models for ALS exist and, if used appropriately, have the potential to improve translation of preclinical approaches and to predict clinical response in ALS. Models should always be selected with respect to the specific biological question being asked. For example, the antisense oligonucleotide against SOD1, tofersen, was assessed preclinically in the SOD1 mutant mouse model, to reflect the relevant human target population for therapy i.e., people with ALS carrying a SOD1 mutation.

In selecting the right model to test a therapeutic hypothesis, diverse approaches are available, including:

- In vitro cell lines e.g. NSC-34 or SH-SY5Y cells, used in screening assays or to establish initial evidence for relevance of a cellular target pathway
- Induced pluripotent stem cells (iPSC) models, which are being increasingly used as tools to explore target engagement and treatment response *in vitro*
- *Ex vivo* autopsy or biopsy samples from people with ALS are critical tools for drug discovery and development, e.g. for the identification of biomarkers and assessment of target expression
- In vivo models including the expression of wild-type human genes, genetically engineered ALS mutations, and models designed to generate specific phenotypes, e.g. conditional (tissue specific) expression to generate neurodegeneration mouse models

Below, we provide some recommendations for how iPSC and *in vivo* models can be used for ALS drug discovery and development.

iPSC models

The role of iPSC models for modelling neurodegenerative diseases has been recently reviewed in detail ⁽³⁶⁾. In ALS drug discovery and development, iPSC-based models may have increasing utility in high throughput and innovative screening approaches ⁽³⁷⁾, and have the potential to facilitate the study of sporadic cases of ALS, which is not currently possible using *in vivo* models ⁽³⁸⁾. For preclinical drug discovery, iPSCs can aid in validation of target engagement, downstream pathway analysis and assessment of the effect of drugs on cellular phenotypes and other readouts which reflect disease pathology. Reprogramming of parent cells to iPSCs may erase the epigenetic signature associated with ageing, potentially reducing the capacity to identify relevant phenotypic ALS markers ⁽³⁹⁾. In contrast, direct reprogramming retains ageing-associated signatures ⁽⁴⁰⁾, and may therefore provide a more realistic tool for validation in specific instances, such as astrocyte toxicity to neurons ⁽⁴¹⁾. Large-scale initiatives to create an accessible bank of iPSC lines from ALS patients are underway, including 'Answer ALS' which is creating up to 1,000 unique stem cell (iPSC) lines from ALS patients and healthy controls ⁽⁴²⁾.

Rigorous and continuous quality control of iPSCs is essential, given the potential for reprogramming to introduce genomic instability ^(43,44) and for culture-driven mutations to propagate though clonal selection. Background

genetic variation, distinct from the disease determining variant, and most of which is not disease relevant, combined with differences in experimental protocols between laboratories, explains why the transcriptomic readouts produced to date show disappointingly little overlap between lines. Isogenic cell lines as experimental controls, and generation of mutant lines on a common parental cell line are approaches being used to limit the confounding effects of genetic background. The iPSC Neurodegenerative Disease Initiative (iNDI) is the largest-ever iPSC genome engineering project, and will model more than 100 mutations associated with Alzheimer's disease and related dementias (ADRD) in isogenic iPSC lines, several of which will be relevant for ALS. Resulting cell lines and phenotypic datasets will be widely shared ⁽⁴⁵⁾.

Recommendations for validation of iPSCs in ALS disease modelling

1. Standardised quality control (46)

- Microbiological sterility mycoplasma, endotoxins, bacteriology, viral testing.
- Genetic fidelity and stability residual vector testing, karyotype, SNP arrays (used in combination).
- Viability e.g. doubling time.
- Identity SNP profiling cells to create cell line "fingerprints" to regularly confirm cell line identity.

2. Comprehensive cell line characterisation

- Initial demonstration that cells retain differentiation potential i.e. ability to form of all three germ layers.
- Characterise cell types against agreed criteria via molecular profiling e.g. neural progenitor cells positive for Pax6+, Sox1+, Nestin+; Hb9+ and Isl1/2+ motor neurons; astrocytes; microglia; muscle ⁽⁴¹⁾.
- Characterisation of cell types via phenotypic hallmarks:
 - Astrocytes: Toxicity to neurons; impaired glutamate uptake (Yamanaka & Komine, 2018).
 - Neurons: Ubiquitination and axonal swelling; Glutamate excitotoxicity; Increased oxidative stress; proteinopathy relevant to the cell model genetic background (TDP-43, SOD1, FUS, C9Orf72 etc.)
 - Microglia: Release of inflammatory mediators ⁽³⁶⁾.
 - Muscle: Fewer muscle unit contractions, increased apoptosis ⁽⁵⁰⁾.
 - Consider use of transcriptomics as a tool to gain more insight into cell identity and biology.

3. Interrogation with positive controls

- Does applying a drug result in the expected change to the phenotype?
- Can the phenotype be modulated either pharmacologically or genetically?

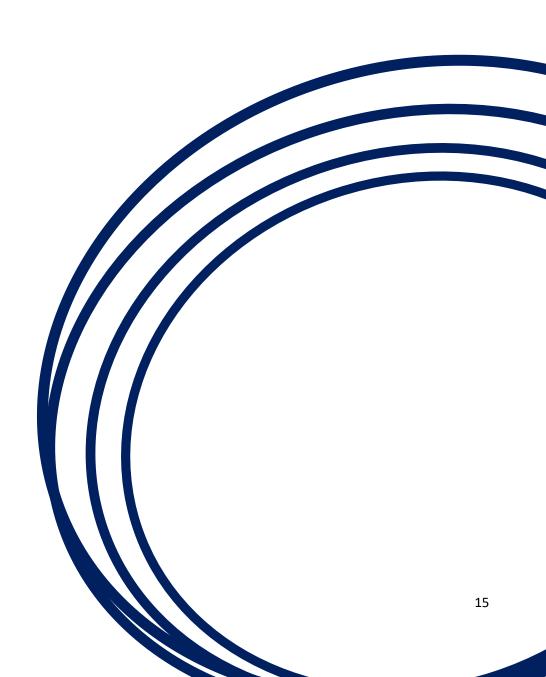
In vivo ALS models

A variety of accessible and well characterized animal models exist for ALS, each of which may have complementary roles in the study of disease mechanisms ⁽⁵¹⁾. However, given that ALS is a disease of mid to late life which may be specific to humans, it is not realistic to expect an *in vivo* model to perfectly recapitulate all features required to function as a precise tool for Go/No-Go decisions about therapy. Individual models should be tailored to address specific questions, and are only one part of a complex decision matrix to take a drug into humans.

In the earlier stages of drug discovery e.g. drug screening, more tractable *in vivo* models that enable high throughput studies include *Caenorhabditis elegans*, *Drosophila melanogaster* and *Danio rerio* (zebrafish). *In vivo* mouse models are key for validation studies in the later stages of preclinical drug discovery and development, for example, for target engagement and PK/PD studies. Mice with disease-relevant phenotypes or pathways can be used to test the effect of drugs on specific aspects of the disease, and to test target engagement. Genetically modified mice expressing ALS related SOD1 mutations, for example, have been critical for the development of

targeted genetic therapies (tofersen). Guidance exists for the use of SOD1 mice in preclinical drug testing ^(7,8). Although, an appropriate model in which to study SOD1 dysregulation, the relevance for most cases of ALS, which is pathologically a TDP-43 proteinopathy, is unclear. Phenotypes in these mice are driven by major overexpression of mutant SOD1 protein, and outcomes based on survival may not translate into human ALS. Moreover, there are no credible mouse models of 'sporadic' ALS, which represents around 90% of all ALS patients, highlighting a key gap in existing models.

Going forward it is critical that investigators use models and assays that are tailored to ask specific questions related to the relevant targets and biology of their preclinical development programme.



Stage in drug	Relevant models examples	Relevant assay examples	Limitations		
discovery/ development					
Pathway screening and analysis	Caenorhabditis elegans Drosophila melanogaster Danio rerio	<i>C. elegans</i> – motility assays <i>D. melanogaster</i> – Motor assays, survival <i>D. rerio</i> – axonal outgrowth, touch evoked escape response, swim assays	 <i>C. elegans</i> and <i>D. melanogaster</i> - evolutionary distant to humans <i>D. rerio</i> – genome duplicated; difficult to manipulate; assays more suited to neuronal development rather than degeneration PK/PD and blood-brain barrier permeability 		
Drug screening			require confirmation in other models e.g., rodents		
D			Metabolism different to humans		
Demonstration of target	Wild-type mouse <u>OR</u>	Biodistribution assays (e.g., PET)			
engagement in vivo	ALS mouse model <u>OR</u>	Tissue and pathway analysis (e.g., spatial			
	Mouse model displaying	transcriptomics, -omics, mass spec)			
	dysregulation of target pathway				
	e.g., non-ALS mouse model of neurodegeneration				
Demonstration of drug efficacy <i>in vivo</i>	ALS mouse model <u>OR</u> Mouse model displaying dysregulation of target pathway	<u>Behavioral assays:</u> Grip strength; motor performance, rotarod; electrophysiology; clinical investigation e.g.,	Electrophysiology is difficult to perform and signal is typically very low during early disease stages		
		overall mobility, weight; behavioral analysis; survival; in vivo axonal transport assays Interventional endpoints:	 Behavioral studies are expensive Clinical investigation in mice is subjective and open to interpretation 		
		Measurement of translational biomarkers; muscle force – isolated muscle preparations; motor unit survival (e.g. neuromuscular junction integrity); axon count	 Interventional endpoints are limited in number of time points as animals are typically deceased upon measurement Variation in technical expertise between laboratories 		

Guiding principles for ALS therapeutic preclinical data packages

To support the development of high-quality preclinical data packages and ensure critical evaluation occurs before assets progress to clinical studies, we recommend the principles below are systematically considered. A robust body of evidence to support and inform well planned Experimental Medicine studies is central to building greater confidence of achieving a positive translational outcome of the right target for the most appropriate patient group and stage of disease progression.

Delineate the biological pathway/mechanism being targeted, and whether its modulation protects cells				
or restores function				
• Studies in relevant <i>in vitro</i> and/or <i>in vivo</i> models, including molecular techniques such as knock-down and overexpression studies.				
Assess evidence that targets are linked to disease and expressed in the relevant tissue in humans				
 Bioinformatic approaches e.g. whole genome sequencing, genome-wide association studies. Target expression studies in relevant human neuropathological tissue e.g. immunohistochemistry, spatial transcriptomics, <i>in situ</i> hybridisation. 				
Identify whether the target is druggable and whether its modulation alters disease progression				
 Assess whether the drug penetrated the target tissue (e.g. CNS, muscle) and achieved sufficient engagement of the target to effect a change. Identify appropriate biomarkers of target engagement and explore downstream pathway analysis through iPSC and <i>in vivo</i> models. 				
Replication of results in an independent lab or across several relevant models can build confidence in the therapeutic hypothesis.				
Identify appropriate biomarkers (new and existing) to measure disease modification				
 Plasma NF-L to evaluate disease progression rate (52). Explore further functional outcomes, -omics and imaging approaches aligned to mechanism of action. 				
Predict accurate human dose				
 Through high quality PK/PD studies, identify drug biodistribution and half life, assess potential for any off target effects that may lead to adverse reactions, and agree an approach for route of administration and dosing in Experimental Medicine and Clinical Studies. 				
Identify possible approaches for patient stratification, and predict appropriate trial duration.				
 Can a subpopulation of patients more likely to respond to the therapy be identified. Examples of stratification approaches include: clinical phenotype, genetics, NF-L levels and the ALSFRS progression rate. 				
 Based on treatment response <i>in vivo</i>, can the length of Experimental Medicine and clinical studies required to detect response to treatment be predicted? E.g., to detect changes in NFL levels or impact on ALSFRS. 				



Replication of preclinical findings in multiple models or through multiple approaches

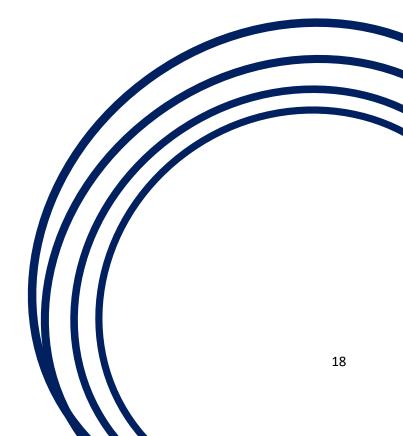
Due to the desperate nature of the disease and the pressure to find effective treatments, ALS therapeutic hypotheses have not always been rigorously validated or replicated before compounds have progressed to human trials, resulting in wasteful allocation of precious resources. Relatively short delays may be incurred at the preclinical stage to invest in building a stronger body of evidence, but the ability to "fail fast" and stop programmes early will speed progress in the long-term.

Embedding a culture where preclinical findings are routinely replicated in multiple models will challenge existing research cultural, funding and infrastructure. Therefore, development of an independent centre or collaborative network that has the appropriate expertise, capability and models to run rigorous preclinical studies could be a valuable resource to the global ALS field.

Phenotypic drug discovery

A counterpoint t to therapy developed against known targets are drugs identified through phenotypic drug screening, an approach with advantages and disadvantages:

Advantages of phenotypic drug screening	Challenges of phenotypic drug screening
Opportunities for drug repurposing e.g. through use of	Labour intensive.
existing compound libraries. Potential for the identification of novel targets.	Difficult to perform at scale, with large compound
Potential for the identification of hovel targets.	libraries.
Utilizes precise readouts reflecting specific aspects of	Target deconvolution is difficult.
ALS pathology e.g. TDP-43 mislocalisation (53),	
modulation of neuronal excitability ⁽⁵⁴⁾ .	



Molecular biomarkers in ALS

Biomarkers for diagnosis, prognosis, PD and monitoring are crucial tools in drug discovery and development, through enabling patient stratification, supporting critical go/no-go decisions, and providing read outs of target engagement and efficacy. In ALS, there has been a lack of well validated biomarkers, which is therefore a major gap in the field which inhibits the development of innovative models for rapidly assessing investigational therapies in patients.

Neurofilaments are key components of the neuronal cytoskeleton. Levels of NFL or phosphorylated neurofilament heavy chain in the cerebrospinal fluid (CSF) rise in response to a range of neurological disorders associated with neuronal loss, broadly in proportion to the rate of disease progression ⁽⁵⁵⁾. Advances in assay sensitivity now permit the reliable measurement of NFL levels in blood, which correlate well with CSF ⁽⁵⁶⁾. Studies in ALS have consistently shown NFL levels to be raised in comparison with healthy and disease control groups ^(56–58)

A role for NFL measurement in the diagnosis of ALS in a specialist setting remains uncertain since it does not increase the diagnostic accuracy of an experienced clinician ⁽⁵⁸⁾. The reasons for delayed presentation of ALS to the neurologist are complex, and whether this can be reduced by using NFL as a screening test for serious neurological disease in a general practice setting is worthy of investigation, since data from UK Biobank suggests that they rise approximately 2 years before symptom onset ⁽⁵⁹⁾. An upward trend in neurofilament level is also found in pre-symptomatic ALS mutation carriers up to two years before transition from asymptomatic to manifest disease ⁽⁶³⁾. Neurofilament levels are strongly correlated with the rate of disability progression in established ALS so have clearer value in prognostic stratification ^(56,61,62). The levels also typically remain stable for the individual patient over their disease course following diagnosis, after an initial rise during the early symptomatic periods ^(56,61–63). The rate of disability progression in ALS is basically stable for individual patients, suggesting that NFL has very significant potential as a pharmacodynamic biomarker in therapeutic trials. In addition, these studies also identified a potential threshold level for neurofilaments levels above which the pathological process appears to become biologically (and clinically) active, laying the ground for the future application of ASO treatment to presymptomatic gene mutation carriers.

In the absence of a highly effective disease-modifying therapy for ALS, evidence for neurofilaments as a marker of treatment response is necessarily indirect. Studies of antisense oligonucleotide therapy in *SOD1*-mediated ALS showed target engagement in the form of reduced SOD1 levels and also a lowering of CSF NFL levels ^(24,68). NFL measurement may have particular value in the triage of candidate drugs in smaller and shorter open-label studies, in order to prioritise those going on to Phase 3 where standard clinical outcome measures are still needed, for example in the EXPERTS-ALS study. We suggest that all ALS studies measure NFL to build the evidence needed to conclusively evaluate its potential as an ALS biomarker. It is likely that studies of at least 6 months duration will be required to show reliable and meaningful changes in NFL.

Ongoing efforts within the ALS research community to identify additional ALS biomarkers have been reviewed recently and include: urinary P75 extracellular domain, urinary neopterin, chitinases, oxidative stress markers, motor unit survival, compound motor action potential (CMAP), miR181 ^(65,69). Better sharing of longitudinal patient samples from well characterised cohorts and systematic collaboration between academic researchers and pharma could be a transformational approach for the field.

Whilst molecular biomarkers remain a focus for future research efforts, many other approaches are being explored in ALS including behavioural, imaging and digital biomarkers. In particular, a focus on the development of digital remote biomarkers for monitoring disease progression at home will enable easier monitoring for people with ALS in a real-world setting.



Precision medicine for ALS

Personalised medicine aims to stratify and select the patients most likely to respond to a given therapy, and is now routine in oncology, where many clinical studies stratify based on molecular profiling. Precision medicine approaches are at an earlier stage in neurodegenerative disorders such as ALS, with tofersen for SOD1 ALS being the only notable ⁽²⁴⁾. Ultimately, in a disorder in which the majority of patients present sporadically, stratification to promote precision medicine will be more complex and include NFL and other biochemical markers, and non-molecular approaches including clinical features, and more complex prediction tools such as the ENCALS model ⁽⁷⁰⁾.

Recent advances in genomic research through large-scale initiatives such as Project MinE ⁽⁷¹⁾ suggest that genetic stratification, including complex polygenic risk scores, will contribute to precision medicine approaches in ALS. Tools such as the Amyotrophic Lateral Sclerosis online Database (ALSoD) have enabled the collation of information about genes and genetic variants connected to ALS, including how they may affect clinical aspects of the disease ⁽⁷²⁾. To maximise the use of data from clinical studies carried out in ALS, the Pooled Resource Open-Access ALS Clinical Trials Database (PRO-ACT) was established ⁽⁹⁾. PRO-ACT provides users with easy access to over 11,600 fully de-identified clinical patient records with more than 10 million longitudinally collected data points. Other initiatives include Answer ALS, which will ultimately generate the largest and most comprehensive collection of ALS data ever amassed and make it freely available to researchers ⁽⁷³⁾. A priority for the field over the coming years will be to integrate data resources such as this to further our understanding of disease heterogeneity and response to treatment. This could lead to the identification of subpopulations of ALS patients who are more likely to respond to a given therapy.

It is estimated that fewer than 5% of people in the UK and 8% in the USA meet the current inclusion criteria for participation in clinical studies ⁽⁷⁴⁾, suggesting that sub-stratification with current trial design is not realistic and could limit the ability to recruit adequate numbers to power high quality clinical studies. An alternative could be to conduct post-hoc analysis of studies to identify responders based on patient subpopulations. However, this requires larger and more costly trials, and sophisticated statistical approaches. Patient stratification may also raise commercial considerations for pharmaceutical companies, where segmenting the population and limiting the label for a subtype of ALS may reduce the incentive to take drugs forward for approval.

Both trial design and patient stratification approaches therefore require urgent attention and will require robust biomarkers, to ensure that the provision of high-quality clinical studies remain feasible for precision medicine in ALS.



The role of Experimental Medicine studies in ALS therapeutic development

'Experimental Medicine' studies, also known as Phase 0 studies in the US, have the potential to reduce late-stage attrition of novel drugs in ALS using futility analysis. These small-cohort studies, positioned between preclinical and phase 1, are performed in patients (i.e., the target population), but are not powered to demonstrate a therapeutic effect or drug safety and tolerance. Instead, they are aimed at providing preliminary information about pharmacokinetics, target engagement, mechanism of action, biomarkers and pharmacodynamics, thus enabling information which is critical to drug development, and may help save time and money that would otherwise have been spent on later phase trials ⁽⁷⁵⁾.

Experimental Medicine studies are common in a number of neurological diseases including Alzheimer's Disease and epilepsy. In ALS, a small number of Experimental Medicine-like studies have been identified, including using flecanide ⁽⁷⁶⁾, thalidomide ⁽⁷⁷⁾ and Triumeq ⁽⁷⁸⁾. It should be noted that these studies are longer than standard for Experimental Medicines studies and include functional endpoints like ALSFRS which would not be usual for a well-designed Experimental Medicine study.

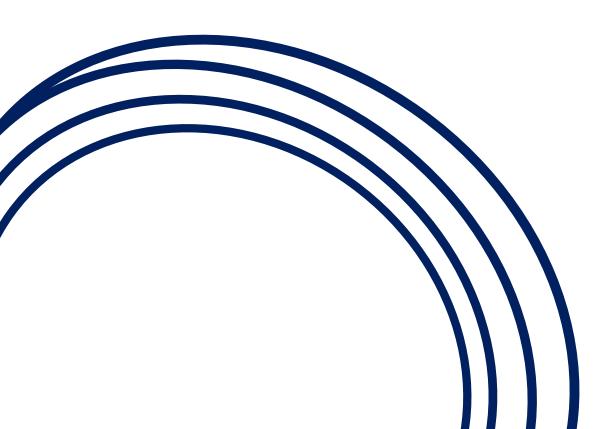
Drug	Rationale	Design	Endpoints	Outcome
Thalidomide	Anti-inflammatory	9-month study	ALSFRS, pulmonary	Negative, adverse side
	(TNF)	of 23 patients	function testing compared	effect profile
			to historical controls,	
			serum cytokines	
Flecainide	Na(+) channel	44-week study	ALSFRS-r,	Safe, "potentially
	blocker and	of 54 patients	neurophysiological index,	effective", not powered
	membrane stabiliser		СМАР	for efficacy
Triumeq	Inhibition of	24-week study	HERV-K expression,	Possible biological
	endogenous	of 40 patients	urinary p75ECD levels,	response supporting the
	retrovirus activity		neurophysiological and	development of a phase
			clinical measures	3 trial.

These studies suggest that an important aim of Experimental Medicine studies is to identify interventions that should not be progressed to later phase clinical study (e.g., thalidomide) in addition to identifying those that show promise.

There are however some key challenges for the successful design of Experimental Medicine studies. Although these aim to be shorter "challenge" studies that provide a more agile response to treatment, it remains essential that study duration is long enough to detect treatment response. Investment in detailed preclinical studies is therefore critical to ensure effort spent on Experimental Medicine studies is productive. Determining the appropriate number of study participants and identifying a relevant patient stratification hypothesis and strategy remains a work in progress. Finally, defining what success looks like will balance the aspiration of 'curative' therapies with the more realistic aim of delivering an incremental benefit in survival.

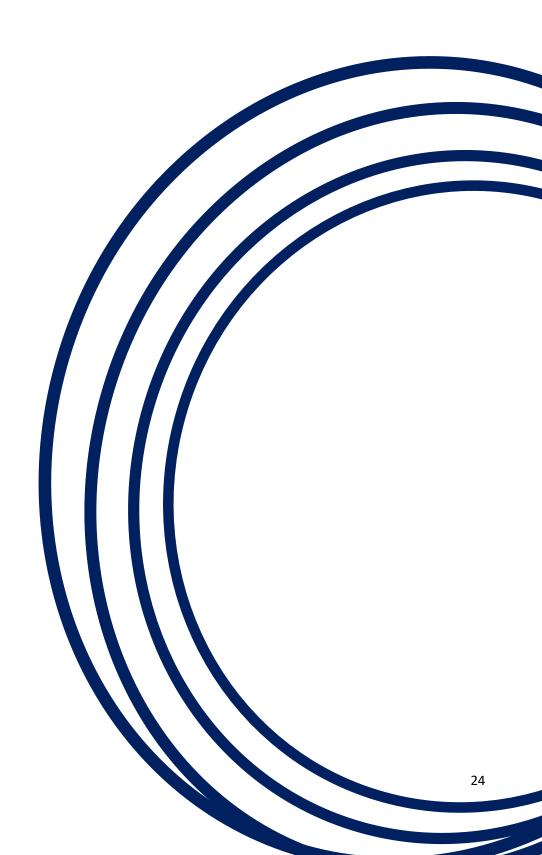
Guiding principles for ALS Experimental Medicine study design and delivery

•	Engage people with ALS in the planning, design and acceptability of the studies.
•	Seek insight on how the results of the study will be communicated to people with ALS.
arge	t engagement and downstream pathway analysis
٠	Assess whether the drug has penetrated the target tissue (e.g. CNS, muscle) and achieved
	appropriate engagement of the target to effect a change.
٠	Determine an accurate dosing strategy for clinical studies.
٠	Demonstrate target engagement through biomarkers, imaging and/or colocalisation studies, or by
	showing modulation of the target pathway or mechanism.
٠	Through exploratory studies such as imaging or -omics analyses, attempt to further understand the
	effect of the drug in cells and tissues of interest, and correlate with disease biology.
oma	arker strategy of treatment response
٠	Assess effect of treatment on NF-L levels and other potential Monitoring Biomarkers identified in
	preclinical studies.
٠	Utilise -omics analyses to identify any potential novel Monitoring Biomarkers.
vide	nce of functional response
٠	Depending on study, target and mechanims of action, and aligned to the preclinical studies conducte
	measure a number of endpoints. This could include Motor Unit Number Estimation of Index
	(MUNE/MUNIX), SVC (slow vital capacity), CMAP, speech, cognitive changes etc. ALSFRS-R may be
	considered, but survival is very unlikely to be included.
•	Determine appropriate treatment duration and clinical study length to observe functional response.
• atier	
• atier	Determine appropriate treatment duration and clinical study length to observe functional response.



Conclusion

The guiding principles outlined here provide an evidence-based framework to inform effective preclinical studies, incorporating lessons from previous drug discovery and development efforts in ALS. Until a successful ALS therapeutic is brought to clinic, we lack a clear exemplar for successful ALS drug discovery and development, but continuous iteration of these principles in the light of new information will improve engagement with key stakeholders, including regulators.



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