Translating molecular mechanisms into ALS risk and patient's well-being

TRΛ

Fondazione Regionale per la Ricerca Biomedica

Sistema Sanitario Regione Lombardia **Giuseppe Lauria**



Amyotrophic Lateral Sclerosis



- Rare disease \rightarrow incidence 3/100,000 inhabitants per year
 - Expected 300 new patient/year in the Lombardia Region
- No disease-modifying treatment \rightarrow dismal outcome
- Sporadic in most; genetic in 15%, increased after NGS availability
- No correlation between phenotype and genotype, even in patients harboring variants in well-established disease-related genes
- Onset and course are variable: impact on care of patients and relatives at risk, and on clinical research
- What determinants could influence onset and course of the disease?













TRANS-ALS Consortium



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- To improve the existing knowledge of disease mechanisms in ALS through an innovative complementary preclinical and clinical approach and to address still unresolved clinical needs.
- To develop new tools to support and monitor research and clinical activities: user-friendly web-based database, genetic and laboratory data for tracking ALS patients and subjects at risk of disease; genetic counseling guidelines; and human and animal repositories of biological samples.
- To positively influence **patients' well-being**
- To make TRANS-ALS Consortium a strategic hub for future national and international academic and industry-sponsored research projects on ALS







- TRANS-ALS adopts a bedside to bench to bedside design, including *in vitro* and *in vivo* functional and validation studies
- Preclinical setting → *in vitro* models to assess the toxic effect of known and newly discovered gene variants on different pathways; *mouse models* with different onset and course on which findings are to be validated.
- Clinical setting → in parallel, to provides deep phenotyping and genotyping of ALS patients and at risk family subjects
 → new hypothesis to be tested in preclinical models







- 1. Signatures of disease onset and progression whose modulation in available *in vitro* and *in vivo* ALS models can prevent or exacerbate disease phenotypes
- 2. New platform of *in vitro* ALS models, including reprogrammed cell models from patients to assess the effect of known and new susceptibility genes identified in ALS patients and to define new druggable targets pathways
- 3. Manipulation of available mouse models to assess the effect of gene silencing, protein toxicity, and other potential susceptibility factors on disease onset and progression
- 4. **Subjects at risk** of disease: in-depth genetic and clinical characterization, to detect prodromal changes and individual risk of disease, to clarify genotype-phenotype correlation, and to investigate the link between immune cell signaling and neurodegeneration







WP1: STUDY OF ALS CAUSATIVE MECHANISMS AND MODIFYING PATHWAYS <u>PARTICIPANTS</u>: IRFMN, UNIMI, CAGRANDA, FMPV-FSM, CNR-IN, FINCB, POLIMI

WP2: SIGNATURES OF ALS ONSET AND PROGRESSION IN PATIENTS AND ADULT AT RISK SUBJECTS

<u>PARTICIPANTS</u>: FINCB, AUX, CAGRANDA, FMPV, AOPGXIII, AOBS, FSM, SPAOLO, S.CARLO, NIGUARDA, AOLEG, OCFM, AOPOMA, AOLEC, AOSAA, HSACOMO, AODV, HUMANITAS

WP3: New tools for ALS Research and Personalized Management

<u>PARTICIPANTS</u>: FINCB, AUX, CAGRANDA, FMPV, AOPGXIII, AOBS, FSM, SPAOLO, S.CARLO, NIGUARDA, AOLEG, OCFM, AOPOMA, AOLEC, AOSAA, HSACOMO, AODV, HUMANITAS

WP4: STRATEGY FOR KNOWLEDGE MANAGEMENT, PROTECTION, EXPLOITATION AND DISSEMINATION

PARTICIPANTS: ALL PARTNERS.

WP5: MANAGEMENT

PARTICIPANTS: ALL PARTNERS

Sistema Sanitario



Results: PRECLINICAL -1



SHARING AND SETTING OF AVAILABLE *IN VITR*O ALS MODELS

- Lymphoblastoid cell line from sporadic and mutated ALS patients. (FMPV)
- VAPB-downregulated motor neuron-like NSC34 cell lines (CNR-IN)
- >50 primary fibroblast cultures from familial (7 SOD1, 16 C9orf72 and 6 TARDBP) and sporadic patients; fibroblast- and blood-derived iPSC (7 C9orf72, 5 TARDBP, 2 sporadic ALS and 8 healthy control) (AUX)
- iPSCs derived from C9ORF72, TDP43, SOD1 patients; iPSCs expressing the DENDRA protein tagged-TDP43 wt or A315T mutation using the CRISPR-Cas9 procedure; iPSCs differentiated in MNs (UNIMI)







STUDY OF ALS ONSET AND PROGRESSION MECHANISMS

- Collection of tissues (blood, CSF, PBMC, brain, spinal cord, nerves, muscle) from fast and slowly progressing mouse models at different disease stages; proteomic analysis in PBMC from ALS patients with different onset and in PBMC and spinal cord (IRFMN & FINCB)
- **Impaired immune response** (MHCI and CD8+T cells) in peripheral nerve of C57SOD1G93A mice (**IRFMN**)
- Increased tRNAs at disease onset in slowly-progressing but not rapidlyprogressing mutant SOD1 ALS mice (neuroprotective stress signaling) (IRFMN)
- Increased mRNA levels (HSPB5, BAG3, p97, p62, Pax7, Myo) in muscles of slowly progressing mice (UNIMI)
- Earlier muscle atrophy in slowly than fast progressing mice at the presymptomatic stage (UNIMI & FINCB)







STUDY OF ALS ONSET AND PROGRESSION MECHANISMS

- Nearly 300 differentially expressed **IncRNAs** found in sporadic patients, whereas a limited amount of IncRNAs deregulated in mutated patients. (FMPV)
- New IncRNA turnover regulatory mechanism involving the binding of the RNA-binding-protein ELAV1/HuR in which the HuR interaction play a stabilization action on InRNAs turnover and exert a regulatory function necessary for the neural stem cells differentiation process. (UNIMI)
- Increase in **VAPB levels** during differentiation of NSC34 cells suggests that this ER-resident protein is important for the elongation of neurites (**CNR-IN**)
- The lack of VAPB induce a delay in neurite outgrowth and an increase in PI4P expression, indicating a critical role in motorneuronal viability (CNR-IN)
- Synthesis and characterization of specific superfluorinated reporters to promote miRNA encapsulation (POLIMI)







C9ORF72-BASED MODELS AND RELATIVE MECHANISMS

- Characterization of the biochemical behavior of 5 different DPRs encoded by RAN translation from the expanded C9ORF72 gene. (UNIMI)
- Confirm that nucleo-cytoplasmic transport is impaired (AUX)
- Differential miRNA expression profile in C9-MNs and in C9-MN-derived extracellular vescicles (UNIMI & FMPV)
- A selective form of **autophagy**, chaperone assisted selective autophagy (CASA) is highly efficient to remove misfolded protein prior to their aggregation (UNIMI)
- The use of ASO in C9orf72 iPSC-MNs reduced the formation of pathological RNA foci by RNA-FISH analysis. (AUX)





Risultati: CLINICA



- Comitati etici
- Biobanca campioni biologici (Mario Negri)
- Banca DNA (FINCB)
- eCRF







	Тіро	Codice ~	Codice Fiscale 0	Data Inserimento	Visite (completate/aperte)	Primo Sperimentatore	Centro Origine o	
0		TRANS-FNB-0001	TRTRSL56A70D761R	01/06/2018	6	Nubilaria Investigator	[FNB] FONDAZIONE IRCCS ISTITUTO NEUROLOGICO CARLO BESTA	Vai al paziente
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0		TRANS-AUX-0001	GRBMRI86L61L219L			Staging Investigator 2	[AUX] ISTITUTO AUXOLOGICO ITALIANO	Vai al paziente
0		RISK-AUX-0001	GGGMMM12P23L325G			Staging Investigator 2	[AUX] ISTITUTO AUXOLOGICO ITALIANO	Vai al paziente

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Legenda:

Risultati: CLINICA



• Protocolli per pazienti

- Diagnosi e presa in carico
- Funzioni fragili
- Neurofisiologia
- Neuroradiologia
- Neuropsicologia
- Disposizioni di fine vita

Protocolli per soggetti a rischio e famiglie

- Percorso presintomatico (valutazioni cliniche, neurofisiologia, neuroradiologia, neuropsicologia)
- Counselling genetico





Risultati: CLINICA



- 120 pazienti (10% mutati, 1 nuova variante)
- 8 famiglie \rightarrow 20 soggetti a rischio





Pubblicazioni scientifiche



PRECLINICA

• 9 lavori pubblicati, 6 sottomessi, 1 in revisione

CLINICA

• 3 lavori pubblicati, 1 in revisione





Meeting del consorzio



- **Kick-off meeting** \rightarrow 15 dicembre 2016
- 1° meeting gruppi preclinici → 24 novembre 2017
- 1° meeting gruppi clinici → 5 maggio 2018
- Meeting gruppo neuropsicologia → 13 giugno 2018

Dicembre 2018 → riunione congiunta risultati intermedi





Ringraziamenti



- Pazienti e famiglie
- Partners del consorzio
- Fondazione Regionale per la Ricerca Biomedica



