

## Experiment Report Form

The double page inside this form is to be filled in by all users or groups of users who have had access to beam time for measurements at the ESRF.

Once completed, the report should be submitted electronically to the User Office via the User Portal:  
<https://www.esrf.fr/misapps/SMISWebClient/protected/welcome.do>

### Deadlines for submission of Experimental Reports

Experimental reports must be submitted within the period of 3 months after the end of the experiment.

### Experiment Report supporting a new proposal (“relevant report”)

If you are submitting a proposal for a new project, or to continue a project for which you have previously been allocated beam time, you must submit a report on each of your previous measurement(s):

- even on those carried out close to the proposal submission deadline (it can be a “*preliminary report*”),
- even for experiments whose scientific area is different from the scientific area of the new proposal,
- carried out on CRG beamlines.

You must then register the report(s) as “relevant report(s)” in the new application form for beam time.

### Deadlines for submitting a report supporting a new proposal

-  1<sup>st</sup> March Proposal Round - **5<sup>th</sup> March**
-  10<sup>th</sup> September Proposal Round - **13<sup>th</sup> September**

The Review Committees reserve the right to reject new proposals from groups who have not reported on the use of beam time allocated previously.

### Reports on experiments relating to long term projects

Proposers awarded beam time for a long term project are required to submit an interim report at the end of each year, irrespective of the number of shifts of beam time they have used.

### Published papers

All users must give proper credit to ESRF staff members and proper mention to ESRF facilities which were essential for the results described in any ensuing publication. Further, they are obliged to send to the Joint ESRF/ ILL library the complete reference and the abstract of all papers appearing in print, and resulting from the use of the ESRF.

Should you wish to make more general comments on the experiment, please note them on the User Evaluation Form, and send both the Report and the Evaluation Form to the User Office.

### Instructions for preparing your Report

- fill in a separate form for each project or series of measurements.
- type your report in English.
- include the experiment number to which the report refers.
- make sure that the text, tables and figures fit into the space available.
- if your work is published or is in press, you may prefer to paste in the abstract, and add full reference details. If the abstract is in a language other than English, please include an English translation.

	<b>Experiment title:</b> Structural basis of regulation of skeletal muscle	<b>Experiment number:</b> LS-2992
<b>Beamline:</b> ID02	<b>Date of experiment:</b> from: 4 <sup>th</sup> October 2021 to: 11 <sup>th</sup> October 2021	<b>Date of report:</b> 3 <sup>rd</sup> March 2022
<b>Shifts: 15</b>	<b>Local contact(s):</b> Peter Boesecke	<i>Received at ESRF:</i>
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**Report: Introduction.** Contraction at the level of the sarcomere, the structural unit of the striated (skeletal and cardiac) muscles is due to the cyclical interactions of myosin motors extending from the thick filaments and the nearby actin containing thin filaments. The aim of this project is to investigate the structural basis of the regulation of the contraction, which involves proteins other than the contractile proteins myosin and actin. In the last five years X-ray diffraction experiments at ID02<sup>[1,2,3]</sup> have demonstrated that, beyond the classical Ca<sup>2+</sup>-dependent thin filament activation that makes the actin available for interaction with the myosin motors, a mechano-sensing mechanism in the thick filament recruits myosin motors from their resting (OFF) state as a function of the load of the contraction. Increasing interest for explaining the molecular mechanism of thick filament regulation focuses on two sarcomeric proteins, (i) the giant protein **titin** and (ii) the **myosin-binding protein C (MyBP-C)**. Titin runs from the midpoint of each thick filament to its tip and then extends from the thick filament tip in parallel with the thin filaments to the end of the sarcomere. Titin forms an I-band spring that likely provides the load to prevent development of inhomogeneity in serially linked half-sarcomeres with different force capability<sup>[4]</sup>, but may also transmit external stress to the thick filament. MyBP-C is bound to the backbone of the thick filament in the central one-third of each half (C-zone) and appears to be involved in the control of the OFF state of the motors<sup>[5]</sup> but also bridges thick and thin filaments with its N-terminal<sup>[6]</sup>. The research is aimed at verifying how Ca<sup>2+</sup>-dependent thin filament activation signal is transmitted to the thick filaments by MyBP-C putative links and at clarifying the role of titin in both the maintenance of sarcomere length homogeneity and thick filament mechano-sensing. For this, we use the intense - highly collimated beam at ID02 to record the low-angle interference X-ray diffraction pattern from intact fibres of frog muscle, reporting the structural changes in the filaments and in the myosin motors. During LS-2721 we have first completed the experiments started with LS-2514, in which rapid changes in force were imposed on the resting fibre to determine the stress sensitivity of the myosin-based reflections in the absence of Ca<sup>2+</sup>. Then, we have established a new protocol to verify if (i) the putative MyBP-C links responsible for the fast communicating path between thin and thick filament and (ii) the putative titin ability to couple the load on the thick filament to the OFF-ON switch of motors are Ca<sup>2+</sup> sensitive. For this it is necessary to record the structural changes in the filaments following a stepwise change in force imposed on the stimulated fibre without the confounding effects of the force and stiffness of myosin cross-bridges. During LS-2721 and, to improve statistics, also part of LS-2791, we have tested the effectiveness of the myosin inhibitor ParaNitroBlebbistatin (PNB)<sup>[7]</sup> in preventing motor attachment and force generation in electrically stimulated fibres. During the last measurement session (LS-2992, Oct 21) the changes in the X-ray signals marking the structural dynamics of the thick filament have been recorded following a 0.25  $T_0$  force step imposed both at rest and in tetanically stimulated fibres in the presence of PNB.

**Muscle fibre preparation and protocols.** Frogs (*Rana esculenta*), cooled to 2-4 °C, were killed by a percussive blow to the head followed by destruction of the spinal cord in accordance with EU official regulations on Use of Laboratory Animals, and of the University of Florence Ethical Committee (in compliance with the rules of the Decreto Legislativo of Italian Government 4 marzo 2014, n. 26). Small bundles of 2-3 fibres were dissected from tibialis anterior muscles, taking care at minimising the length of the tendon attachment at the two ends. The bundles were then transferred to an experimental chamber containing Ringer's solution at 4 °C and mounted vertically at beamline ID02 between a capacitance force transducer and a loudspeaker-coil motor, carried by micromanipulators for adjustment of the bundle length and position in the X-ray beam path. Two mylar windows were moved as close as possible to the preparation to reduce the X-ray path in water. X-ray patterns were collected on the FReLoN CCD detector with 2048 x 2048 pixels (active area 50x50 mm<sup>2</sup>). The isometric tetanic force at 2.15  $\mu\text{m}$  ( $T_0$ ) was preliminarily measured to calibrate, for each fibre, the size of the force step (0.25  $T_0$ ) imposed either at rest or during tetanic stimulation with the contraction inhibited by the presence of PNB (20  $\mu\text{M}$ ). The sarcomere length (SL) was then set to 2.6  $\mu\text{m}$ , to prevent that at rest the lengthening response following a 0.25  $T_0$  force step could encompass the range of movement of the motor-length transducer. The X-ray patterns were recorded with a camera length of 1.6 m (to collect up to the sixth order of the myosin-based reflections). Radiation damage was minimised by translating the bundle along its axis by 100  $\mu\text{m}$  between X-ray exposures and using fast tandem shutters to limit the exposure to the acquisition time. To follow the stress dependent structural dynamics of the thick filament, we collected 3 ms frames at various times following the force step. Corresponding 3 ms frames in the time series from each protocol were added to improve S:N.

**Results.** Data analysis has been delayed by contingent problems of re-calibration of the FReLoN CCD detector but provisionally reveals an activation dependent enhancement in the ability of titin to transmit the stress to the thick filament and a hierarchy in the ensuing sequence of structural changes concerning thick filament backbone, myosin motors and MyBP-C related signals.

**Conclusions.** The results show that the fastest structural change in the thick filament following the force step is the plastic-like extension (ca 2.5 % per  $T_0$ , time frame centered at ~5 ms after the step) that is similar at rest as during tetanic stimulation<sup>[8,9]</sup>. In the active fibre the stress imposed on the half-sarcomere is transmitted to the thick filament with an I band titin lengthening that is one order of magnitude smaller than in the resting fibre, in agreement with the activation-dependent increase in titin stiffness reported in purely mechanical works<sup>[4]</sup>.

**References.** 1. Linari *et al.* 2015, *Nature* **528**:276-279. 2. Reconditi *et al.* 2017 *PNAS*.**114**:3240-45. 3. Piazzesi *et al.* 2018 *Front. Physiol.* **9**:736-743. 4. Powers *et al.* 2020, *J Physiol* **598**:331-45. 5. Reconditi *et al.* 2014 *J Physiol* **592**:1119-37. 6. Luther *et al.* 2011, *PNAS* **108**:11423-28. 7. Kovacs *et al.* 2004, *J Biol Chem* **279**: 35557-63. 8. Squarci *et al.* 2021, bioRxiv <https://doi.org/10.1101/2021.08.06.455239>. 9. Squarci *et al.* 2022, *Biophys J* **121**:515a

