Director of Centre of New Technologies of the University of Warsaw, with the Project Leader, announce opening of the competition for the position of PhD Student in the Biomolecular Modelling Laboratory – Centre of New Technologies of the University of Warsaw.

**JOB OFFER**

<table>
<thead>
<tr>
<th>Position in the project:</th>
<th>PhD Student</th>
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<tbody>
<tr>
<td>Laboratory:</td>
<td>Biomolecular Modelling Laboratory</td>
</tr>
<tr>
<td>Scientific discipline:</td>
<td>Physical sciences/chemical sciences/biological sciences</td>
</tr>
<tr>
<td>Keywords:</td>
<td>protein structure, biomolecular evolution, molecular dynamics simulations</td>
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<tr>
<td>Job type (employment contract/stipend/other):</td>
<td>stipend</td>
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<tr>
<td>Part-time/full-time:</td>
<td>full time</td>
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<tr>
<td>Number of job offers:</td>
<td>1</td>
</tr>
<tr>
<td>Stipend amount/month:</td>
<td>5000 PLN gross</td>
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<tr>
<td>Position starts on:</td>
<td>01.06.2024 or as soon as possible afterwards</td>
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<tr>
<td>Maximum period of contract/stipend agreement:</td>
<td>26 months, with possible extension to 36 months</td>
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<tr>
<td>Institution:</td>
<td>Centre of New Technologies, University of Warsaw</td>
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<tr>
<td>Project leader:</td>
<td>Dr. Piotr Setny</td>
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<tr>
<td>Project title:</td>
<td>From ancestral peptides to contemporary proteins: folding and stability of protein building blocks</td>
</tr>
<tr>
<td>Competition type:</td>
<td>Sonata Bis 10</td>
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<tr>
<td>Financing institution:</td>
<td>National Science Centre</td>
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</tbody>
</table>

Contemporary protein structures most likely evolved by merger of short peptides capable of adopting simple yet stable structural forms. Such basic protein building blocks, so called supersecondary motifs, are typically formed by two regular secondary structure elements linked by loops. Bioinformatic analysis indeed reveals a number of such conserved, ancient units in otherwise unrelated protein domains. Given the idea that they survived during evolution, likely acting as folding and stability seeds for proteins, we will investigate their properties using computational analysis and molecular dynamics simulations. Specifically, our goal will be to find out if they possess any extra stability in comparison to other, structurally similar but sequentially diverse supersecondary motifs, and if yes, what physical properties contribute to this stability. Answering those questions will shed light on: a) the mechanism by which proteins fold overcome
unfavourable hydration contributions, b) the emergence of hydrophobic protein cores, c) the principles allowing the design of new protein structures.

**Key responsibilities include:**

- setting up and conducting molecular dynamics simulation
- developing custom analysis methods
- preparing scientific presentations and manuscript drafts
- participating in seminars and conferences

**Profile of candidates/requirements:**

**Practical requirements:**

- basic knowledge of Linux operating system and scripting language (e.g. Python, Perl)
- good knowledge of English
- previous experience with molecular modelling would be welcome, although it is not strictly necessary

**Formal requirements:**

- The competition is open for persons who meet the conditions specified in the regulations on the allocation of resources for the implementation of tasks financed by the National Science Centre Sonata Bis 10 grant.
- MSc degree in physics, chemistry, biology or related discipline. The MSc degree should be obtained before the date of starting work in the project.
- Confirmed status of a PhD student in Poland (on the date of starting work in the project at the latest).

**Required documents:**

1. Cover letter
2. Current curriculum vitae with the list of scientific activities undertaken to date
3. Contact details of the former supervisor or a recommendation letter sent directly to the project leader (p.setny@cent.uw.edu.pl)
4. Copy of MSc certificate (or, if the MSc certificate has not been obtained yet, a certificate/document about the date of MSc defense)
5. Document confirming the status of PhD Student (to be provided before starting work in the project);
6. Signed information on the personal data processing.

**We offer:**

- work in a friendly atmosphere, in a modern and well equipped institute
- Extensive training in computational modelling of complex biomolecular systems
- Opportunity to participate in courses and conferences
- Access to state of the art computing equipment

**Please submit the following documents to:**  
[link to email]

**Application deadline:**  
30.04.2024

**Date of announcing the results:**  
10.05.2024

**Method of notification about the results:**  
Email, website: [link to website]