

Services offered by EU-OPENSREEN

Service/Tool Name	Description	Contact	Mode of access
Assay adaptation	<p>Once a robust, sensitive and physiologically relevant bench-top assay is in place, dedicated assay adaptation groups, in close collaboration with the User, will adapt bench-top protocols into an High-throughput screening (HTS) format.</p> <p>This assay adaptation process will involve a strong emphasis on quality, including: quality control of reagents; (patho-)physiological relevance; pharmacological consistency; robustness; minimised variability; and tracking of statistical descriptors (e.g. signal to background and the Z' factor).</p> <p>A pilot library 5,000 compounds is available to help validate assays and potentiate larger projects.</p>	office@eu-openscreen.eu	Users are requested to submit project proposal to the EU-OPENSREEN office
Compound/drug screening	<p>HTS of an assay against the EU-OPENSREEN chemical collection (> 100,000 compounds); <i>in silico</i> profiling, including basic chemoinformatic analysis and identification of frequent hitters and other potential false positives; hit selection; confirmatory screening, including orthogonal assay and IC/EC50 determination; basic counter screening; basic SAR based on screening data; QC of confirmed hits.</p> <p>Screening platforms provide state-of-the-art technologies, e.g. absorbance, luminescence, fluorescence, (time-resolved) FRET/BRET, FLIPR, and AlphaScreen, and implement assay formats such as cell-based, biochemical and (to a limited extend) model organisms-based assays.</p>	office@eu-openscreen.eu	<p>Users are requested to submit project proposal to the EU-OPENSREEN office.</p> <p>A compound replenishment fee structure applies to all projects and users from countries which are members of the ERIC will receive a discount on this cost.</p>

<p>(Medicinal) Chemistry services</p>	<p>Evolution of structure-activity-relationships (SAR) to enable the design of new improved compounds. Depending on the optimization aim, a variable set of biophysical and pharmacological profiling data as required for the project will be collected (e.g. compound solubility in aqueous solutions, cellular membrane permeability (PAMPA, Caco-2), stability in liver microsomes (mouse, human), serum protein binding, cytotoxicity), and, if requested, pharmacokinetic evaluation in mice or rats, and <i>in vivo</i> general tox (zebrafish, mouse).</p>	<p>office@eu-openscreen.eu</p>	<p>Users are requested to submit project proposal to the EU-OPENSSCREEN office</p>
<p>Bioprofiling of donated compounds</p>	<p>All compounds donated by the user which enter the EU-OPENSSCREEN ERIC compound collection are characterised and annotated for basic physico-chemical (e.g. identity, solubility, light absorbance and fluorescence) and essential to know biological properties (cytotoxicity, antibiotic, antifungal etc.) by testing in a standard panel of assays.</p>	<p>office@eu-openscreen.eu</p>	<p>Users are requested to contact the EU-OPENSSCREEN office</p>
<p>Provision of standardised data</p>	<p>All data generated through QC/bioprofiling and screening activities will be published in EU-OPENSSCREEN's open-access database with an optional 'grace' period (i.e. delayed publication of data). Large-scale open-access data provide the basis for computational data integration to obtain a systematic view, allow for prediction of drug-target interactions and networks as well as of adverse effects and drug combinations.</p>		<p>Open-access. (The access to the database will utilize the facilities of ChEMBL.)</p>
<p>Training and education</p>	<p>Training of second (master) & third (PhD) cycle students, postdoctoral scientists and principal investigators in chemical biology in compound management, process automation, IT, assay development and screening.</p>		