



Compounds for HTS  
Chemical building blocks  
Fragment libraries  
Targeted libraries  
Drug discovery services

# **GABA RECEPTORS ANTAGONIST LIBRARY**

OTAVA offers GABA Antagonist Library. This library provides an excellent basis for drug discovery projects related with GABA receptors.

The library consists of **862 compounds\***.

## All compounds are:

- **in stock**; available amounts: 1 – 50 mg
- **Drug-like only**; reactive, pan-assay interference (PAINS), redox-active and aggregator compounds were removed from the library.

## QA/QC passed:

- minimal purity of compounds is **90%**;
- by **NMR** and/or **GC/LC/MS**
- **NMR spectra are available** upon request

## Friendly packing services:

- **Cherry-picking is available**
- Supplied as dry powder or DMSO solution\*\*
- Packaging in deep-well plates or barcoded vials\*\*\*
- **Weighing out is free**

\*Please note that the library does not contain known inhibitors. The compounds were selected with computational approach and are intended for screening projects

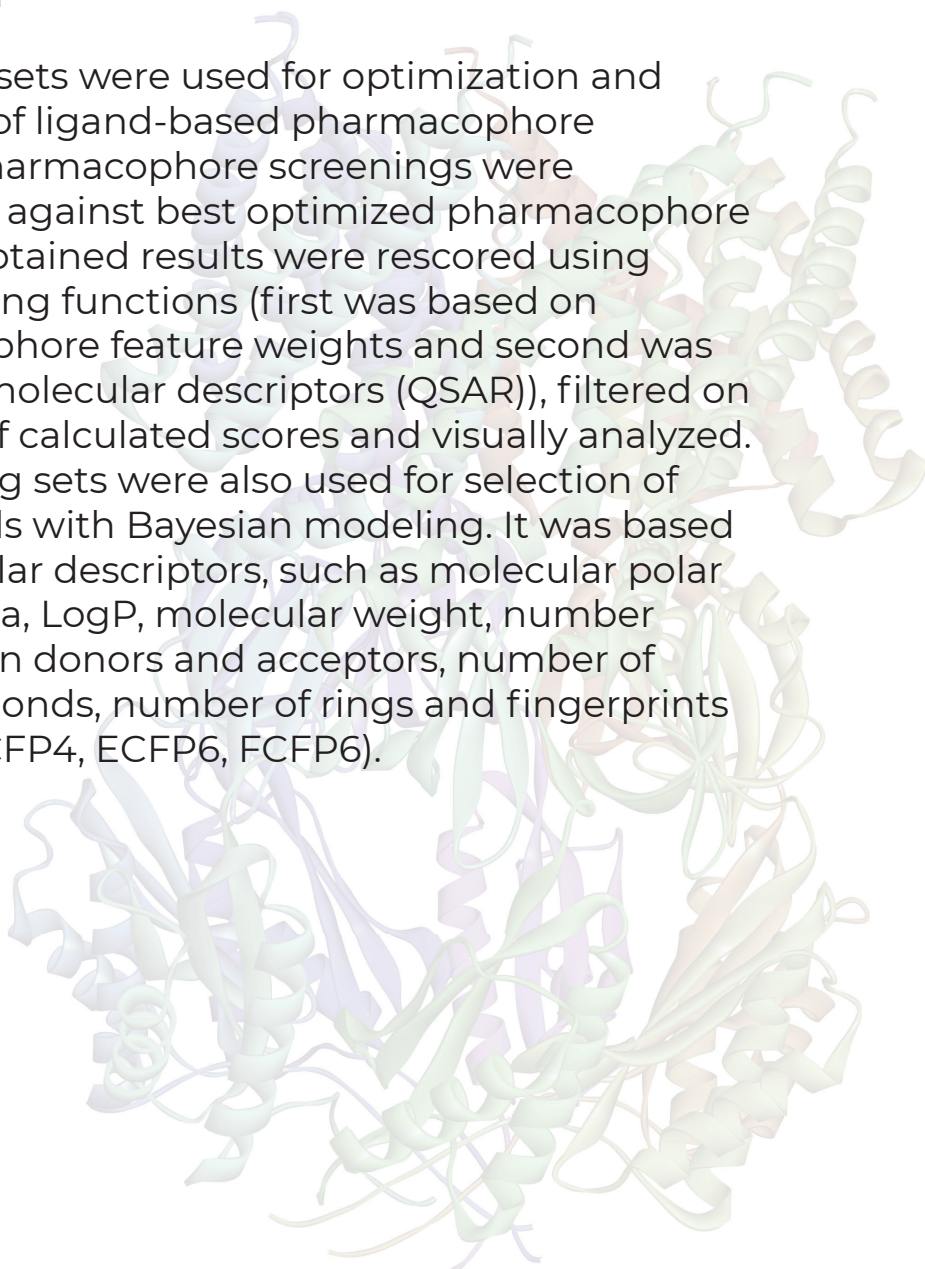
\*\*there is additional fee for preparation of the solution

\*\*\*4 ml amber glass vials or Deep-well plates: Matrix cat# 4247 (1.4 mL, Blank, Polypropylene, Round Bottom Tubes) w/CapMats. Or plates and vials provided by customer.

## Design speciality:

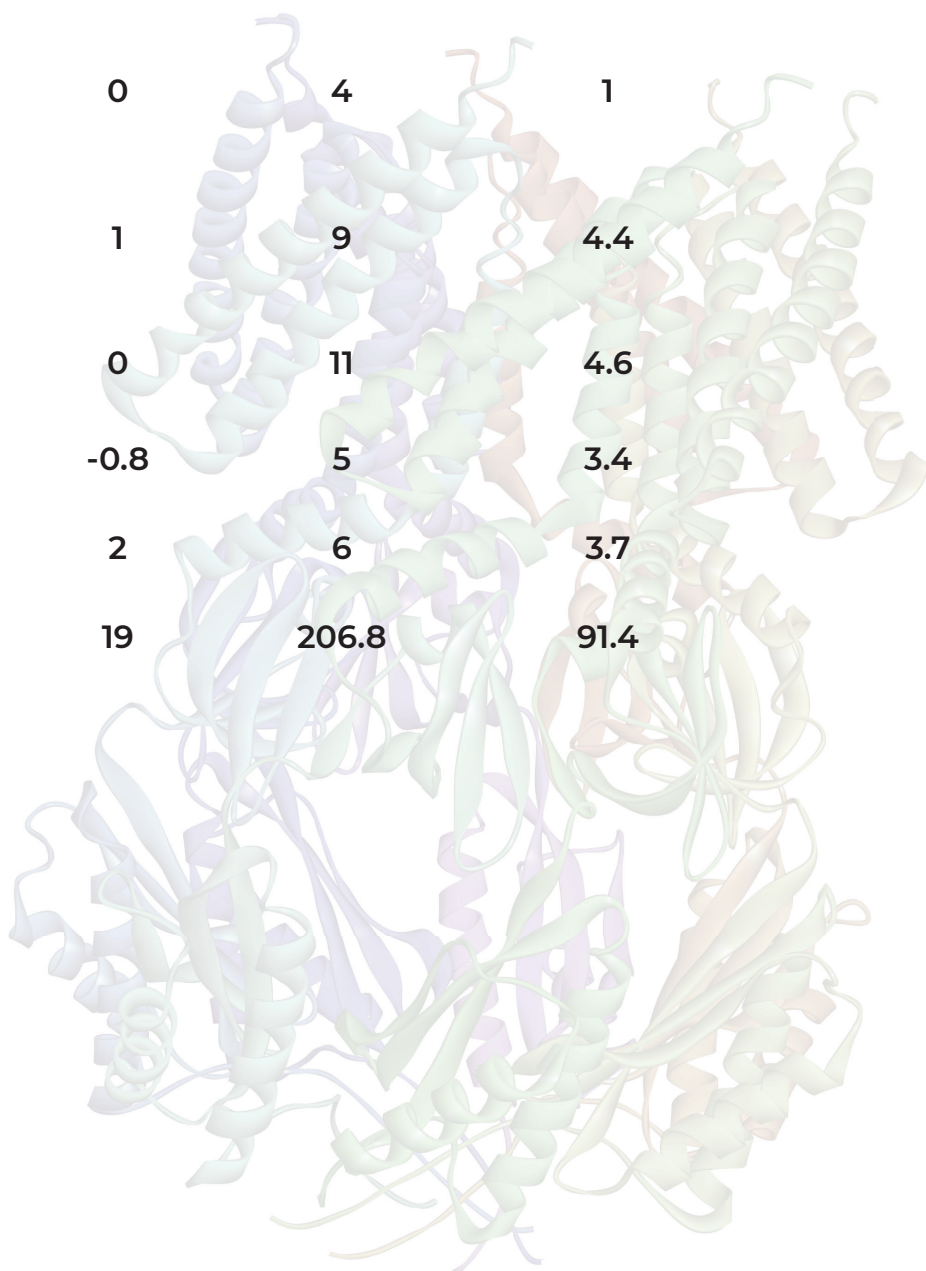
These special sets of compounds have been carefully designed with powerful predictive tools - Bayesian and pharmacophore modeling combined with QSAR (See Appendix 1). Antagonist of GABA were taken from ChEMBL database. Corresponding training sets were formed based on these compounds.

Training sets were used for optimization and validation of ligand-based pharmacophore models. Pharmacophore screenings were performed against best optimized pharmacophore models. Obtained results were rescored using two rescoring functions (first was based on pharmacophore feature weights and second was based on molecular descriptors (QSAR)), filtered on the basis of calculated scores and visually analyzed. The training sets were also used for selection of compounds with Bayesian modeling. It was based on molecular descriptors, such as molecular polar surface area, LogP, molecular weight, number of hydrogen donors and acceptors, number of rotatable bonds, number of rings and fingerprints (ECFP4, FCFP4, ECFP6, FCFP6).

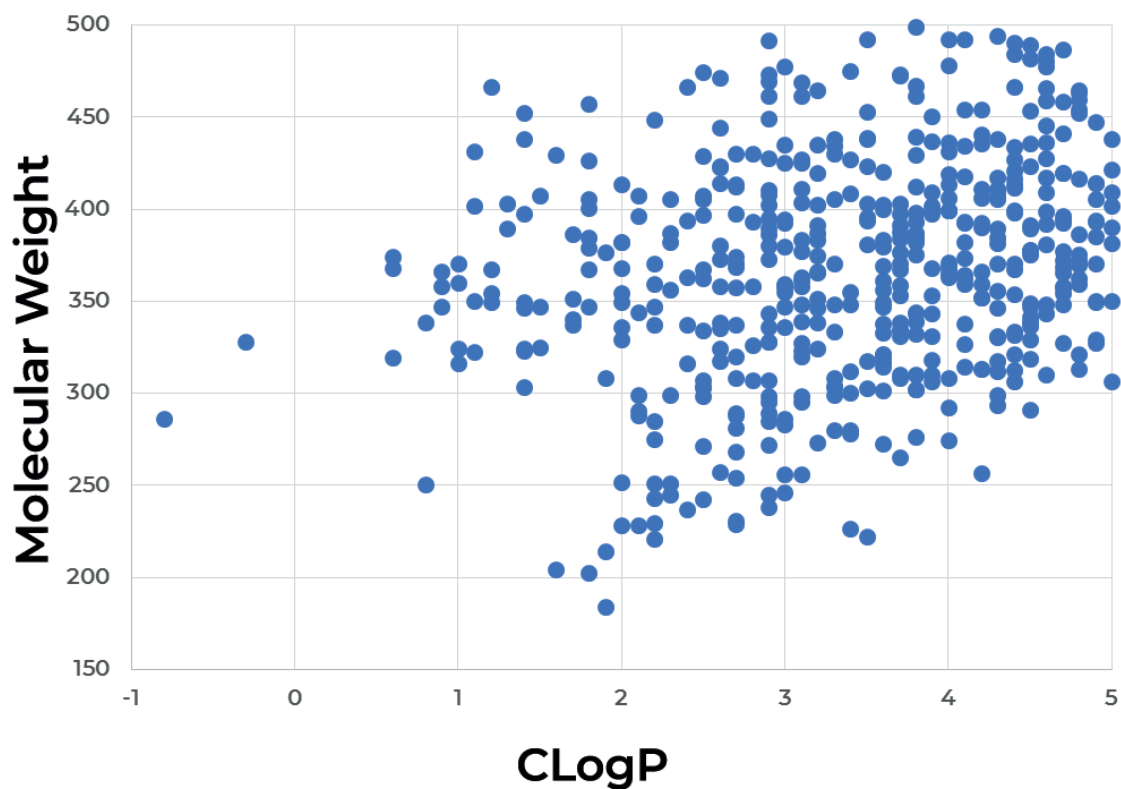
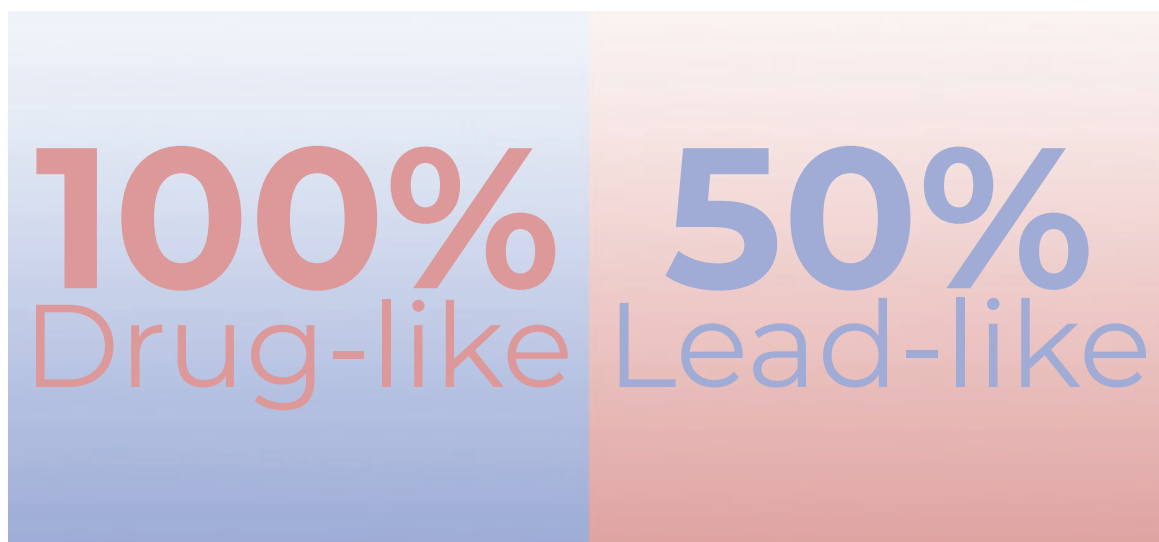


## The summary of the library characteristics:

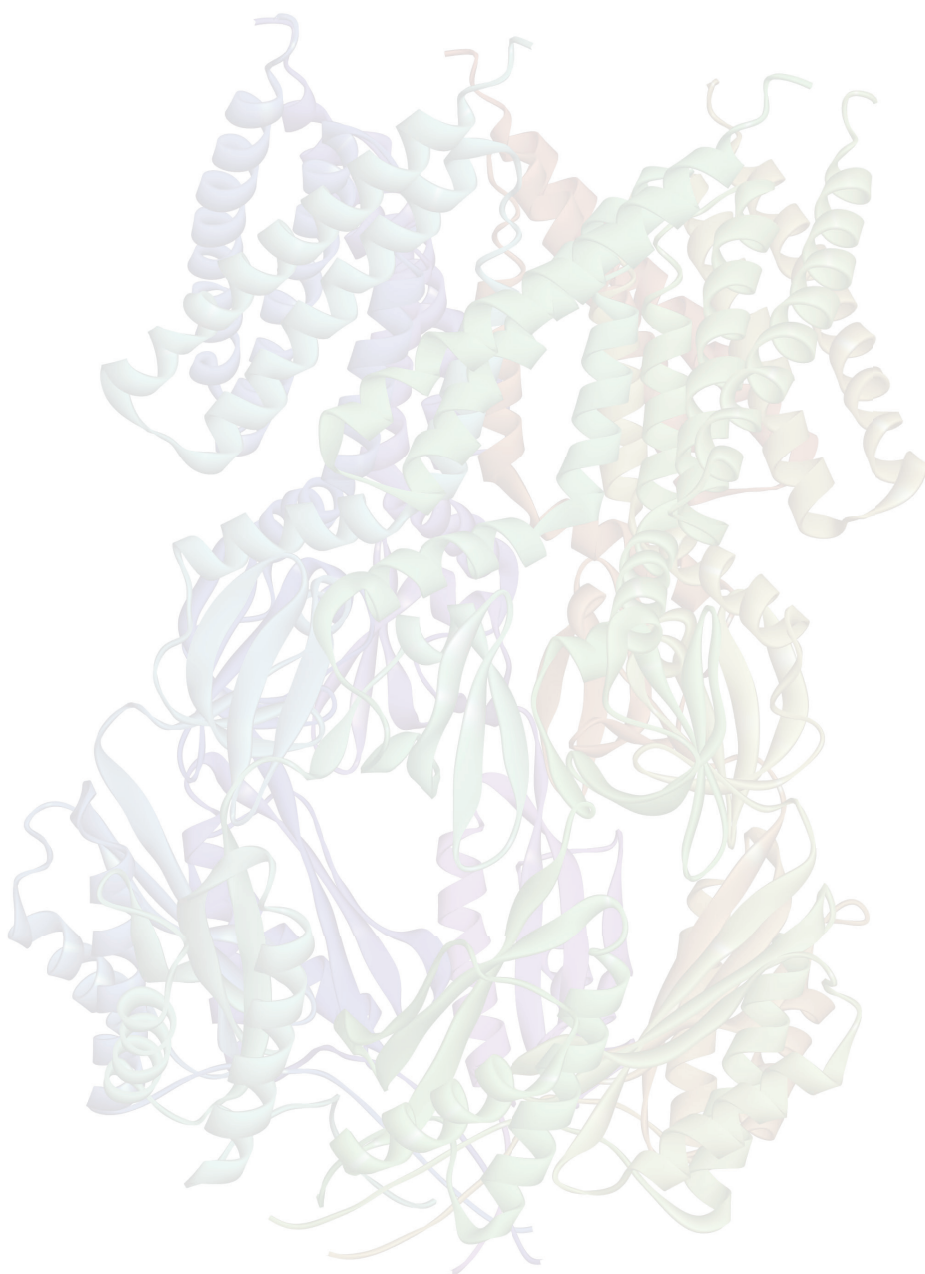
	Minimum	Maximum	Average value
<b>Molecular Weight</b>	184.2	500.6	367.4
<b>Number of Hydrogen Bond Donors</b>	0	4	1
<b>Number of Hydrogen Bond Acceptors</b>	1	9	4.4
<b>Number of Rotatable Bonds</b>	0	11	4.6
<b>CLogP</b>	-0.8	5	3.4
<b>Number of Rings</b>	2	6	3.7
<b>Polar Surface Area</b>	19	206.8	91.4



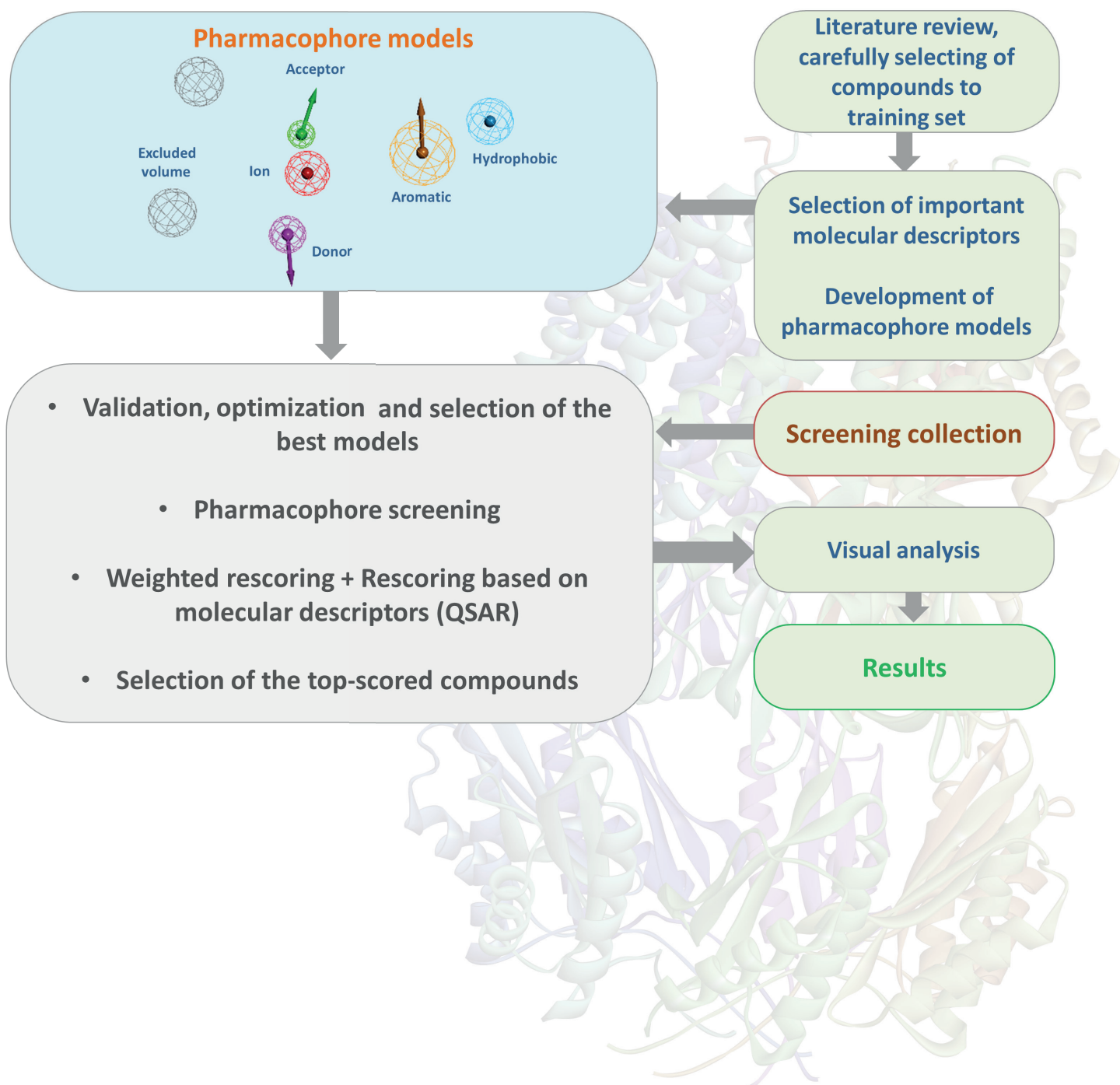
# Distribution of physicochemical properties of compounds in the library:



## Appendix 1



## Scheme 1. Application of ligand-based pharmacophore modeling for targeted library:



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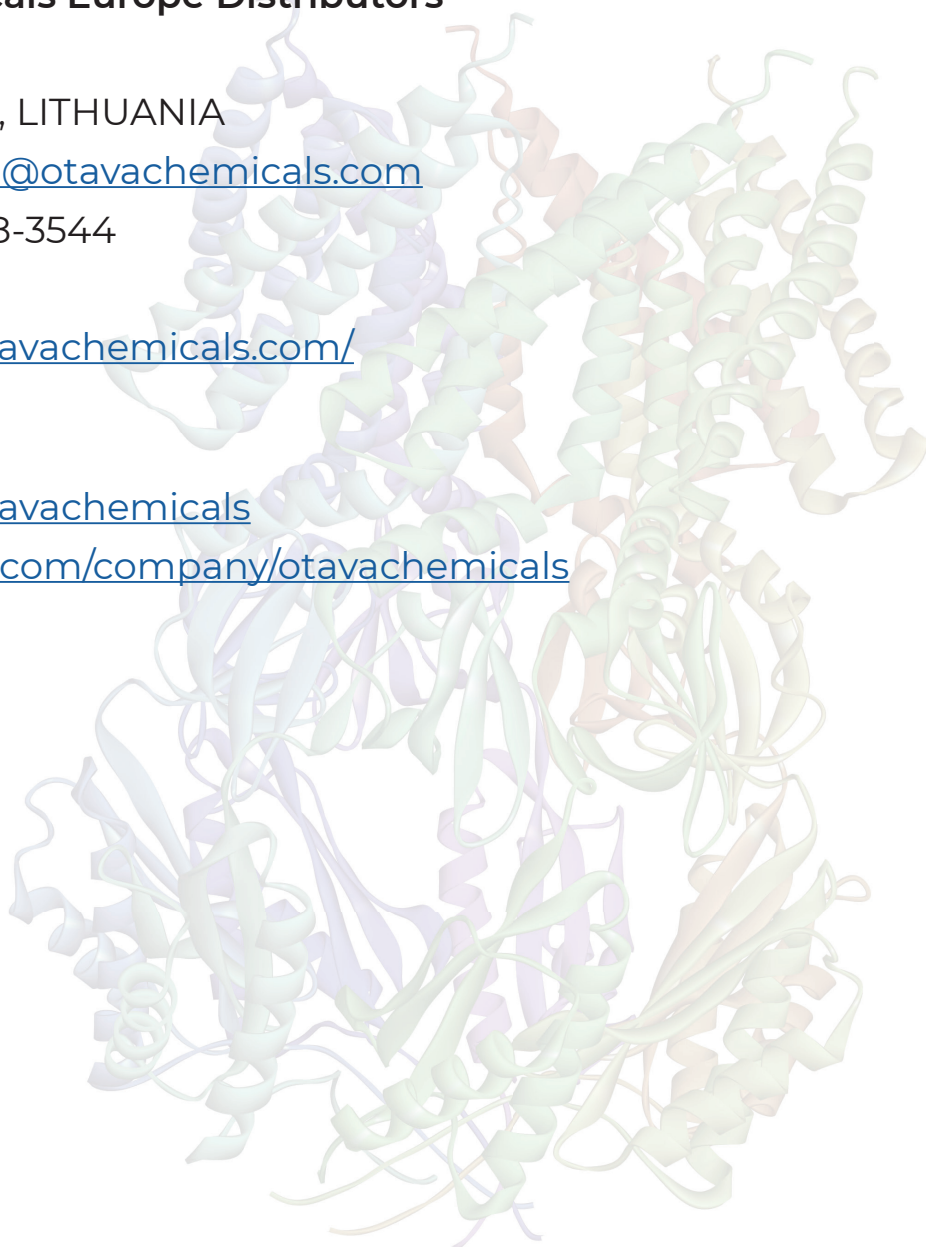
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