

Concentration-response modeling of in vitro bioactivity data from complex mixtures of priority Superfund compounds

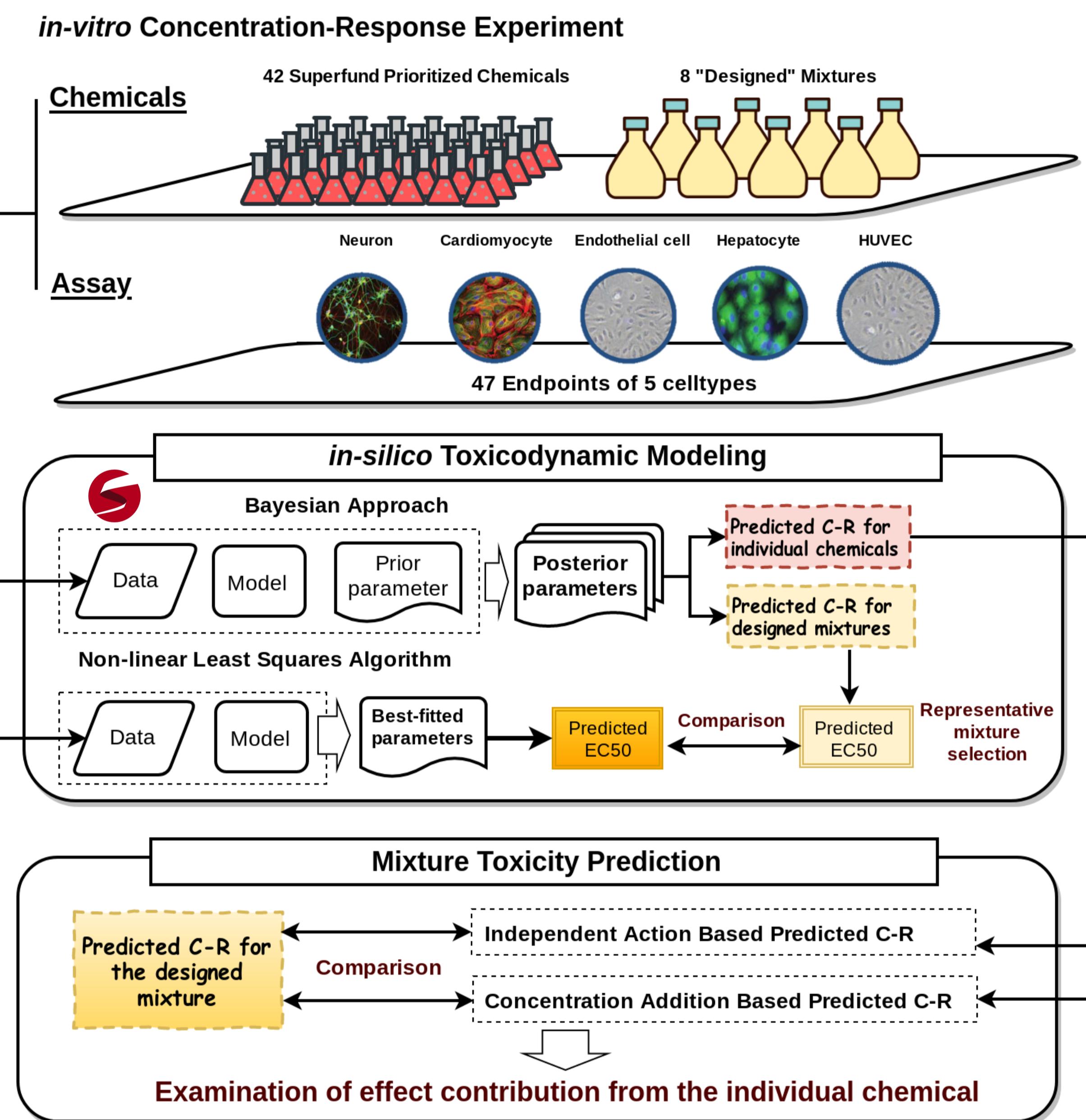
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INTRODUCTION

- Environmental chemicals at Superfund sites are composed of diverse compounds that include heavy metals, pesticides, industrial chemicals, polycyclic aromatic hydrocarbons, and plasticizers.
- Traditionally, most exposure-effect studies focus on the adverse effects of a single chemical or a mixture with few compounds. The approach might not reflect the "real-world" exposure scenario that contains dozen of pollutants and can cause additive or synergistic effects of human health.
- The high throughput screening-based toxicity testing with *in-silico* approach provides the opportunity to more fully examine the biological responses from complex mixtures.

WORKFLOW



METHODS

- At first, the sigmoidal dose-response model that derived from Fréchet distribution function (also known as Weibull distribution) was used to quantify concentration-response profiles from individual chemicals and mixture [1].
- The traditional Marquardt-Levenberg non-linear least-squares (NLS) algorithm [2], as well as Bayesian approaches [3], were used. Two algorithms were compared to determine the better method to describe the concentration-response properties [4].
- Making the predictions as to mixture toxicity under the assumptions of independent action (IA) and concentration addition (CA), and compared the predictions with mixture concentration-response data [5].
- Finally, we investigated the effect of contributions from the individual chemicals in the designed mixture.

The study objects include:

- To examine the robustness of the probabilistic approach in quantifying the bioassay concentration-response profile.
- To address the property of toxicity from the designed mixture that was composed of dozens of chemicals.
- To identify the contribution of bioactivity responses from individual chemicals and conduct the prioritization.

