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TITLE: Machine Learning Based Modelling of Human and Insect Olfaction Screens Millions of compounds to Identify Pleasant Smelling Insect Repellents

AUTHORS: Joel Kowalewski[1], Sean M. Boyle[2], Ryan Arvidson[2], Jadrian Ejercito[2], and Anandasankar Ray[1,2]

AFFILIATION:

[1]Interdepartmental Neuroscience Program

[2]Department of Molecular, Cell and Systems Biology

University of California, Riverside, CA 92521, USA

CORRESPONDING AUTHOR: Anandasankar Ray

Department of Molecular Cell and Systems Biology, University of California Riverside, 3401 Watkins Drive, Riverside, CA 92521, USA, Phone:+1(951) 827-5998, Email: anand.ray@ucr.edu

Abstract

The rational discovery of behaviorally active odorants is impeded by limited information on how the olfactory system generates percept or valence for a volatile chemical. In previous studies, we showed that chemical informatics could be used to predict ligands for a large repertoire of odorant receptors in *Drosophila* (Boyle et al., 2013). However, it remained difficult to predict behavioral valence of volatiles since the activities of a large ensembles of odor receptors encode odor information, and little is known about the complex information processing circuitry. This is a systems-level challenge well-suited for Machine-learning approaches which we have used to model olfaction in two organisms with completely unrelated olfactory receptor proteins: humans (~400 GPCRs) and insects (~100 ion-channels). We use chemical structure-based Machine Learning models for prediction of valence in insects and for 146 human odor characters. Using these predictive models, we evaluate a vast chemical space of >10 million compounds *in silico.* Validations of human and insect behaviors yield very high success rates. The discovery of desirable fragrances for humans that are highly repulsive to insects offers a powerful integrated approach to discover new insect repellents.

Introduction

Animals can detect volatile molecules in their environment using specialized protein receptors in their olfactory organs. Mammalian odorant receptors were discovered in 1991 and are G-protein coupled receptors (GPCRs) numbering >400 in humans (Buck L and Axel, 1991). Several years later starting in 1999 three families of distinct insect odorant receptors were discovered from the *Drosophila* genome which are ligand-gated ion channels and number ~100 (Clyne et al., 1999, 2000). Over the subsequent 20-30 years, studies in mice and insects focusing on odor detection, odorant receptors, odorant receptor neurons, and higher-level circuitry, have revealed important principles. A single volatile is usually detected

by a combination of receptors, and the activity from the neural circuit processing the information is

extremely complex. Previously, we successfully developed a chemical informatics approach to discover

ligands for a large number of odorant receptors in *Drosophila* (Boyle et al., 2013). However, the

mechanisms connecting olfactory perception (e.g., rose, lemon, pleasant, unpleasant, etc.) to olfactory

receptor activity are incompletely understood and it is therefore challenging to predict behavior from

odorant receptor activity alone. There is also often a lack of correspondence between simple, 2D

representations of odorants and their perceptual descriptions, and the molecular-cellular organization of

the olfactory system lacks a simple discernible coding strategy that enables predictions, unlike auditory

or visual processing (Buck L and Axel, 1991; Grabe & Sachse, 2018; P Mombaerts, 1999; Peter Mombaerts,

2001; Sachse & Beshel, 2016; Vassar et al., 1993).

Yet Machine Learning algorithms have shown promise in accurately predicting human odor

perception (Keller et al., 2017; Kowalewski et al., 2021; Kowalewski & Ray, 2020b; Lee et al., 2023;

Sanchez-Lengeling et al., 2019). Recently, we have shown that the volatile structure-to-percept prediction

can be improved by using the activities of a small number of human odorant receptors and applying

machine learning algorithms to identify highly predictive combinations (Kowalewski & Ray, 2020b). In

insects, we have limited success modeling the behavioral valence of volatiles from olfactory receptor

neuron activity in the fruit fly, *Drosophila melanogaster*. Since receptor activity depends on chemical

structure, this suggests machine learning algorithms can potentially detect a consistent physicochemical

basis for simple behavioral or complex perceptual responses (Kowalewski & Ray, 2020b; MacWilliam et

al., 2018).

To test whether Machine learning based modeling can lead to identification of strong behaviorally

active volatiles, we selected a challenge that has a very high potential for easing human suffering and yet

had little innovation: pleasant smelling fragrances that are also repulsive to harmful insects that transmit

diseases to humans or cause agricultural loss. Annually, mosquitos account for ~500,000-700,000 deaths

and a billion infections due to vector-borne illnesses like malaria, Dengue, Yellow Fever. Insects also

destroy ~30% of the global agricultural produce causing additional suffering. Insects are currently

managed through widespread application of insecticides, but mutations are emerging that confer

insecticide resistance and affect insect olfactory responses in both human disease vectors and plant pests

(Deletre et al., 2019; Duneau et al., 2018).

The most widely used topical repellent for mosquitoes is DEET (N,N-Diethyl-m-toluamide), a

chemical with undesirable chemical and cosmetic properties. DEET (N,N-Diethyl-m-toluamide) was

discovered >75 years ago through empirical testing. It is not widely used outside the developed world,

where vector borne diseases are present globally, and are also use restrictions due to its strong solvent

properties, even dissolving plastics and nylons. Recent studies have also raised human health concerns ,

as the chemical interferes with inhibitory neurotransmission in mammals via $GABA_A$ and Glycine receptors

(Grant & Hall, 2019). Unfortunately, not enough is understood about the odorant receptors for aversion

in mosquitoes as a result there has been limited innovation in the search for repellents even a quarter

century after discovery of insect odorant receptors. The last repellent registered by the US EPA for

commercial use was Picaridin, over 20 years ago.

Volatile compounds that have pleasant fragrances are desired by humans in cosmetics and are

therefore excellent candidates for novel mosquito repellents. In order to identify fragrances that would

be aversive to insects, we assumed machine learning (ML) could be used to identify structural features

enriched among repellents and then detect these features within any arbitrary, large chemical space

(Maldonado et al., 2006). We trained a support vector machine (SVM) on a comprehensive set of

chemicals with known repellency from literature and identified predictive physicochemical features. In

parallel, we adapted a collection of recently developed ML models to predict 146 different human odor

percepts, helping us organize the candidate repellents by their potential fragrances (Kowalewski et al.,

2021; Kowalewski & Ray, 2020b). We then validated the accuracy of behavioral predictions for insects and

humans by randomly testing 50 predicted chemicals for repellency in mosquitoes and odor perception in

humans. The validation success was >75% in each assay, repellent and perceptual, with ~95% of the

chemicals tested for repellency in round 1 showing aversion, which left ~5% as neutral or slightly attractive

to mosquitoes. Collectively, this suggests a strong physicochemical basis for odor valence and percept in

insects and humans. In addition, we also built *in silico* toxicity models, filtering the compounds across

multiple parameters for human use. A small number of the fragrances were tested on evolutionarily

distinct insects, finding that they are repellent to mosquitoes and the fruit fly, *Drosophila*.

Tens of millions of dollars and several years may be required for identification and adequate

human-safety analysis for registration of new repellent chemicals by the EPA (Gupta & Bhattacharjee,

2006). Machine learning approaches for modeling toxicity as well as other physical and cosmetic attributes

are steadily advancing, suggesting it is possible to accelerate this search and significantly reduce costs.

Accordingly, we included additional computational filters to select predicted repellents such as predicted

fragrant odor perceptual qualities. Because fragrant molecules that are also insect repellents may be rare

to find, we applied this approach to a large library (>10 million) of purchasable chemicals from the ZINC

15 database. These efforts represent a dramatic expansion of the insect repellent chemical space and

provides a theory-based high-throughput method to drive rational discovery of novel insect repellents.

Results

Machine learning can learn to predict aversive valence in insects from chemical structure

To predict insect behavior, we developed a novel chemical informatics approach using Machine Learning

(ML). We assembled a training set of volatile chemicals known in the literature to have a negative

behavioral valence including eucalyptol, linalool, geraniol, citronella, alpha-thujone, beta-thujone (Kline

et al., 2003; Klocke et al., 1987; Syed & Leal, 2008), DEET, picaridin, 34 N-acyl piperidines that are

structurally related to picaridin (Katritzky et al., 2008) and structurally diverse odorants from fruits and

food sources that are not expected to elicit aversion (Carey et al., 2010; Hallem & Carlson, 2006). We

generated energy-minimized 3D structures for each and calculated 3,242 physicochemical features for

201 training chemicals. From it we identified 18 features that were highly correlated with aversive valence

(correlation of 0.912). We then fit a support vector machine (SVM) with these features to learn to predict

the repellency of test chemicals (Figure 1).

The trained model successfully predicted the valence of the test chemicals (avg AUC = 0.994),

indicating that it could distinguish volatiles with aversive valence from non-repellents (Figure 1).

Evaluation of the predictive chemical features indicated six membered rings, carbon-nitrogen distances,

tertiary amides, and the positioning of oxygen atoms in the molecule are associated with repellency

(Figure 1).

We subsequently used this repellency model to evaluate a >440,000 small molecules assembled

from databases which includes ~3,000 chemicals from natural sources like plants, insects, vertebrates,

and compounds that are in food, beverages, and cosmetics. The top 500 predicted compounds (top ~0.1%

of hits) represent a structurally diverse group of chemicals, with some showing similarities to volatiles in

the training set (Figure 1). Several naturally occurring compounds were also in the list of precited

repellents. Since the ML model successfully classified known repellents versus non-repellents, we next

experimentally tested the candidate repellents.

Machine learning can prioritize candidate repellents by modeling their odor qualities to humans

Importantly, repellents must meet many physical and cosmetic demands for human use. However,

evaluating the scent characteristics of candidate repellents with human subjects is a laborious, expensive,

and time-consuming task. In order to create a method to bypass this bottleneck, we adapted our recent

perceptual modeling approach involving the prediction of 146 odor perceptual descriptors from chemical

structure; example perceptual descriptors include "fruity other than citrus", "lemon", "orange",

"cinnamon", "perfumery" and unpleasant ones such as "sickening", "rancid", "animal", and "dirty linen"

to name a few (Kowalewski et al., 2021; Kowalewski & Ray, 2020b). The models were trained with data

from prior human behavior studies in which participants evaluated ~150 odorants and mixtures according

to the 146 descriptors. The development of the methods and validation of the models were rigorously

evaluated in the previous publications (Kowalewski et al., 2021; Kowalewski & Ray, 2020b). Using these

models, we predicted odor descriptors for the candidate insect repellents that were discovered using the

model described in Figure 1. The compounds had a wide variety of predicted odor characteristics covering

both pleasant and unpleasant fragrance profiles. Together with the insect valence predictions, this gives

us the foundations for selecting desirable fragrances that can protect humans and plants from pests.

Validation of computational behavioral predictions of human smell and insect aversion

To validate the behavioral predictions of scent or repellency, on humans and insects, respectively,

we selected 44 compounds based on their availability and affordability. First, we compared them to

reports from human testing of smell properties. While odor qualities remain poorly characterized, 37 of

the 44 tested had odor and taste annotation in a sensorial databases at the The Good Scents Company.

We could then rigorously evaluate the accuracy of the trained perceptual models by ROC analysis. The

perceptual models predicted the observed descriptors for most volatiles with a high success rate (Avg.

AUC = 0.77) (Figure 2A,B). Interestingly, many chemicals in the predicted repellents have pleasant odor

characteristics, including highly desirable ones such as fruity, floral or perfume-like, such as seen in the

word cloud of enriched predicted characters for the predicted insect repellents (Figure 2C).

We next tested the compounds against *Aedes aegypti* mosquitoes. We developed a high

throughput assay utilizing the innate attraction of the female mosquitoes to a heat block approximating

human skin temperature (37 °C) covered with a filter paper (Figure 3A). Each of the test compounds were

individually evaluated by applying on the filter paper (3% in acetone) and counting the numbers of

mosquitoes landing on the surface with solvent control before and after applying the compound. A

repellency index indicated that most of the predicted compounds (~90%) showed >50% repellency to

female *A. aegypti* mosquitoes (Figure 3B). These results suggest that the machine learning (ML) algorithm

was capable of predicting a chemical's perceptual qualities for humans and its valence for insects. This set

of chemicals represents a very large increase in new insect repellents, comprising diverse chemical

structures and smell characteristics that are desirable for human use.   In a separate study we had tested

a few of these repellents using a human arm-in-glove behavioral assay and observed strong repellency

(Boyle et al., 2016), suggesting that the newly discovered compounds are likely to exhibit similar effects.

Since we do not know the molecular receptor targets for these diverse repellents, and different

insect species have both highly diverse and conserved olfactory receptors, we cannot tell *a priori* whether

the repellency is narrowly effective in mosquitoes or conserved across diverse insects. To determine

broad-spectrum aversive valence among insects, we tested another distantly related dipteran, *Drosophila*

melanogaster, which is >250 MY apart evolutionarily. All of the predicted compounds displayed

concentration-dependent aversive valence to *Drosophila* in a trap-based 2-choice assay (Reeder et al.,

2009; Syed et al., 2011). Most of the tested compounds showed strong repellency (Figure 3C). The

simplest interpretation is that the machine learning algorithm was able to identify chemical features that

predict negative valence across multiple species of dipteran insects.

Repellents show calcium mobilization in mammalian cells

An alternative approach to identifying insect repellents would be to computationally model a known

cellular or molecular activity relevant to repellency. Interestingly, one of the aversive compounds used in

the repellency training set is the commercial repellent DEET, had previously been seen to cause calcium

mobilization in human HEK293 cells at high concentrations (Dennis et al., 2018). In order to further

evaluate if such Ca^{2+} mobilization is a common feature for repellency, we performed additional

experiments.

When HEK293 cells were exposed to DEET, a dose-dependent calcium transient was observed,

with an EC50 of 56.6 mM (~1% vol/vol), below the concentrations used on skin applied formulations (7-

98% vol/vol) of which up to 60% is absorbed into the skin (Schoenig et al., 1996; Selim et al., 1995). (Figure

4A). Additional cell lines from different organisms were tested, including an insect S2 cell line, and all of

them were sensitive to DEET at similar EC50s (Figure 4A). The DEET-induced calcium transients were

similar in magnitude to the positive control, a 100 µM ATP induced signal (e.g., GPCR mediated) (Figure

4B). To assess the source of DEET induced calcium flux, cells were depleted of intercellular calcium with

treatment with thapsigargin (a SERCA antagonist) 10 minutes before the assay. These calcium-depleted

cells did not show increased Ca transients when treated with DEET. This indicates that DEET is likely

inducing release from intercellular calcium stores, in a dose-dependent manner (Figure 4C). Interestingly,

two additional commercially used repellents, picaridin and diethyl benzamide (a structural analog of

DEET), also evoked a strong intracellular Ca^{2+} signal (Figure 4D). Because these three compounds were

part of the training data to build our machine learning insect repellent models, we tested whether the

newly identified aversive fragrances also showed similar activity. We tested a sampling of the verified

aversive fragrances, and a few traditionally known repellents, at two concentrations (5% and 0.5%) in the

HEK293 cell line and found that there were indeed some that showed Ca^{2+} release (Figure 4E). However,

there appeared to be no correlation between the repellency index and the Ca^{2+} activation of the fragrant

repellents (R^2=0.0093), suggesting that this conserved effect is unlikely to contribute to aversive behavior.

Nevertheless, the data provided an opportunity to develop a better understanding of the

physicochemical basis of the Ca^{2+} mobilization seen *in vitro* and we trained machine learning models to

predict the calcium activity (Figure S1A). Upon analyses, the top chemical features for predicting the

calcium activity were different from the features selected by the model to predict repellency. This too

suggest that while there is a structural basis for basis for calcium activity which does not overlap with

shared features for detection as a repellent. The features fell into 3 broad categories including 2D

autocorrelation, RDF, and 3D MoRSE features. Autocorrelations are distance-based metrics, with the 2D

autocorrelation referring to the number of bonds between a pair of atoms in the molecule, while radial

distribution function (RDF) features encode dependencies between atoms that give rise to molecules'

characteristic infrared spectra (Hemmer et al., 1999). The 3D MoRSE features are based on theoretical

electron diffraction patterns and are therefore sensitive to differences in 3D geometries (Devinyak et al.,

2014). In previous work, these features have been shown to be predictive of ligand bonding and biological

activity (Beglari et al., 2020; Cheng et al., 2010; González et al., 2006).

Evaluation of models trained with these features on test data showed a high correlation between

the predicted and observed Ca^{2+} activities (Figure S1). Using another approach, where we classified the

chemicals as broadly falling into one of two categories (high versus low activity), the predictions were also

accurate, further emphasizing that the Ca^{2+} signal does indeed have a strong physicochemical basis.

Collectively, this suggests the unique 3D shape, rather than generic, possibly indiscriminate information,

underlies the observed Ca^{2+} activity, supporting our contention that the activity may be a persistent off

target effect among certain chemical repellents and should therefore be further investigated. While the

physiological consequences of this effect have not been studied, we incorporated this activity as a

parameter into our machine learning pipeline, allowing us to further triage repellents for future use by

low estimated Ca^{2+} activity, in addition to other desirable characteristics for human or animal use.

Mining massive commercially available chemical spaces for novel insect repellents

The size and diversity of the chemical library we initially screened (~0.45M) is in the range of high-

throughput wet lab screening that pesticide discovery companies can undertake. The Machine learning

approach allows us to screen far more without the limitations of cost of synthesis and screening time. To

demonstrate the massive throughput capacity of this approach, we increased our search space to >10

million commercially available chemicals (from the ZINC 15 database), canvasing far more structural

diversity than the ~450,000 chemicals we previously screened. We built a new training set including the

chemical repellents we experimentally verified, more than doubling the number of the actives while

adding greater structural diversity. The top features fell mostly into 2D and 3D autocorrelations and 3D

MoRSE categories, suggesting that certain types of features became increasingly important following the

addition of more repellent activities. Notably, some features changed from this vastly improved training

set; that is, moving away from simple features that account for amides, or more generally, chemical bonds

including nitrogen (N). This possibly reflects an initial bias toward DEET, an amide, that was resolved when

including more complex feature selection methods and increasing the size and diversity of the training

set.

There was a strong correlation between the predicted and observed repellency when validating

these updated models on test data (R = 0.73). Then, by applying the filters that we progressively

incorporated into the pipeline, and described throughout (e.g., insect valence, odor perceptual qualities,

and intracellular calcium release), we identified a set of candidates still numbering in the hundreds. We

next predicted the acute oral mammalian toxicity (Rat) using a computational model from our earlier work

(Kowalewski & Ray, 2020a) and eliminated all compounds falling into EPA Tox categories I and II. The top

predictions contain many aminobenzoates comprised of benzene, amino, and ester groups. Some

chemicals notably differ in that they resemble amide repellents like diethyltoluamide (DEET) (Figure 5A).

The top hits were predicted to have a variety of different human odor characters, many of which are

perceptually pleasant or are useful in agriculture (Figure 5B). This approach accelerates the identification

of novel, fragrance-based, broad spectrum insect repellents.

Discussion

Here, we have developed a comprehensive machine learning pipeline to screen small or large

chemical spaces for suitable alternatives and verified numerous predicted chemicals for repellency in

various behavioral assays and across species, indicating the effectiveness of the pipeline. The rapid

identification of fragrances that have excellent behavioral activity, pleasant smell and safety greatly

improves the likelihood of regulatory approval and potential for use.

Interestingly, we identified a strong intracellular Ca^{2+} signal in response to DEET and picaridin as

well as some anthranilates. While the mechanism remains unclear, it has been previously observed in

cultured cell lines after DEET exposure (Dennis et al., 2018; Swale et al., 2014). One possibility proposed

in the literature is that the elevated intracellular Ca^{2+} is due to activation of octopamine receptors.

Octopamine plays a modulatory role, with targets including OctαR and OctαβR in insects and TAAR1 in

mammals (Kleinau et al., 2011; Swale et al., 2014), and while this could explain the many complex

physiological responses that have reported in the DEET literature (Briassoulis, 2001; Clem et al., 1993;

Corbel et al., 2009; Tavares et al., 2019), an unknown but conserved receptor subtype is still possible.

Additional research will be needed to determine whether this phenomenon is important for off-target

mammalian toxicity. In either case, the model we developed to predict the Ca^{2+} mobilization may help

isolate a set of key physicochemical features, enabling identification of compounds that have this effect.

However, since the calcium activity does not explain the repellency, additional electrophysiological and

genetic experiments are needed to further clarify the mechanism of action.

Although many repellents have been studied, DEET remains the most widely used commercially

available chemical repellent. Yet research continues to emerge about off-target toxicities in humans who

are repeatedly exposed to DEET; specifically, accounts suggest adverse skin reactions and activities in

human or broadly mammalian nervous systems (Briassoulis, 2001; Grant & Hall, 2019; Osimitz et al.,

2010). This study demonstrates the successful development and application of a machine learning

pipeline that accelerates research into insect repellency and its physicochemical basis. The methods and

data presented provide a high-throughput and rational way to identify additional novel repellent

chemicals. Here, we emphasized screening candidate chemicals that are most likely to fit multiple

requirements (smell and repellency) rather than simply repellency. As these requirements steadily

increase, it is obvious that massive, unexplored chemical spaces offer the most promising leads. By

predicting and filtering 10+ million purchasable chemicals according to their desirability for human use,

we further illustrate an essential role for computational approaches (e.g., machine learning) in future

repellency studies, accelerating the discovery of safe, ecologically friendly, and effective repellent

chemicals.

Material and Methods

Behavior testing

Repellency Assay in Drosophila

Repellency was tested using a *Drosophila melanogaster* Canton S strain (RRID:BDSC_1), backcrossed >5

generations to w1118 (RRID:BDSC_3605) strain in 2-choice trap assay. All chemicals were purchased at

highest purity from Sigma Aldrich (RRID:SCR_008988), TCI, Alfa Aesar, Penta, Parachem, Bedoukian, and

Molbase. A chamber was built from DRAM vials, with four air holes made in the plastic lid and the bottom

of chamber is coated with 5 ml of 1% agarose to maintain moisture. Traps are made from 1.5 ml Natural

Microcentrifuge tubes with bottom 3 mm cut to form an opening and a 1000 ul blue graduated pipette

tip catalogue number 1111-2023. To create a plastic funnel, a pipette tip was cut on both ends; cut to 1.0

cm from tip end to expand the opening of the funnel and then cut on the other end of plastic to create a

final height of 2.5 cm for funnel. A BioRad PCR 0.2 ml Tube cap was cut from the strip and was inserted

into the lid of microcentrifuge tube. Test odor (35 ul) was placed in the PCR tube cap and lure (90 ul) was

placed in outer ring between the PCR cap and the microcentrifuge cap. In the dark, 4–7-day old, starved

flies (10 males, 10 females) were released per trap and followed for 2 days. Counts were taken manually,

and the Preference Index (PI) was computed using the formula below.

$$Preference\ Index\ (PI) = \frac{(\#\ flies\ in\ treated\ trap) - (\#\ flies\ in\ control\ trap)}{(\#\ files\ in\ test\ trap) + (\#\ flies\ in\ control\ trap)}$$

Repellency Assay in Mosquitoes

A non-contact heat attraction assay was performed by simultaneously offering mated female *Aedes*

aegypti mosquitoes that were ~4-10 days old, a test chemical on a filter paper coupled with an attractive

heat source. Heat for the assay was provided by a Hot hands® Hand warmers HH2 (Heat Max, Dalton, GA)

which was activated by shaking in gloved hands. Temperature (~37°C) was verified using IR thermometers.

The hand warmer was fitted snugly into a 100 mm x 15 mm petri dish base and covered with 15 cm x 15

cm piece of polyester net secured round the petri dish by a pair or 8-inch plastic cable ties (Gardner

Bender, Milwaukee, WI). The petri dish with hand warmer was covered with a larger 150 mm X 15 mm

petri dish with a 10 cm x 7.5 cm rectangular window cut out. This netted window aligned with the hand

warmers exposing a heated surface.

A polyester net (measuring 9 cm x 8 cm) was treated with 500 µl of either acetone (solvent) or 3%

(v/v or w/v) test chemical and allowed to air dry for 30 minutes to evaporate solvent. The test compound

($\sim$0.2mg/cm^2) treated net was secured by stretching between a frame made using two 10 cm x 7.5 cm

flexible magnetic frames with 7.5 cm x 6.2 cm cut out windows. This dose is 88% lower than the amount

recommended in the WHO, WHOPES "Guidelines for Efficacy Testing of Mosquito Repellents for Human

Skin" recommended dose ($\sim$1.67mg/cm^2) actives. Three more flexible magnets were added on top as

spacers from the heat source. The net-magnet assembly was placed directly over the 10 cm x 7.5 cm cut

out openings of the larger 150 mm X 15 mm petri dishes holding the heat pads. The stimulus was placed

on the top net of the frames. Temperature was monitored by a laser Infra-red thermometer and assay

started at 37°C. A puff of 5% CO_2 was blown into the cage to behaviorally activate the mosquitoes and

initiate the assay. A webcam timelapse is used to record mosquitoes landing on the net for automated

counting offline using ImageJ (RRID:SCR_003070). A custom Macro is used in ImageJ (RRID:SCR_003070)

to count the mosquitoes in the selected area (the net) across the timepoints. The counts from each frame

are automatically tabulated.

Macro:

run("8-bit");

setAutoThreshold("Default");

//run("Threshold...");

```
328        setThreshold(0, 80);

329        run("Analyze Particles...", "size=2-70 show=[Overlay Outlines] clear summarize stack");

330         dir = getDirectory("image");

331          name = getTitle;

332          index = lastIndexOf(name, ".");

333          if (index!=-1) name = substring(name, 0, index);

334          name = name + ".csv";

335          selectWindow("Summary of "+getTitle);

336          saveAs("Measurements", dir+name);
```

Since this is a 1-choice assay, for each cage and compound, a solvent alone control is done before and

after the test material, with at least 20 minutes between the trials. Any significant changes in behavior

between the control solvent should indicate if mosquito behavior is acceptable.

Repellency coefficient = (mean cumulative number of mosquitoes on solvent treated control netting –

mean cumulative number of mosquitoes on treated netting)/ mean cumulative number mosquitoes

attracted to solvent treated control netting. Cumulative number of mosquitoes on the window of

treatment are counted for at time points 2,3,4,5 min. Each compound has 2-23 trials, ~40 mosquitoes per

trial.

In vitro assay of Ca^{2+} mobilization

All mammalian cell lines were maintained at 37 C, 5% CO2, and 99 percent relative humidity. Insect

Schneider 2 (S2) Cells (RRID: CVCL_Z232**)** were maintained at 26 C and 99% relative humidity. HEK293T

(ATCC, RRID:CVCL_0063), CHOk1 (RRID: CVCL_E8IB) and NG108-15 (RRID: CVCL_0464) were grown in

DMEM media (Gibco), supplemented with 10% FBS (Gibco), and antibiotic-antimycotic supplement

(Gibco). S2 cells (RRID: CVCL_Z232**)** were cultured in Shields and Sang M3 Insect Medium (Sigma),

supplemented with Insect Medium Supplement (Sigma). Cells were transfected with the luminescent

calcium reporter aequorin 24 hours before experiment with X-tremeGENE 9 transfection reagent, in the

case of HEK293T (and CHOK1, and X-tremeGENE HP, in the case of NG108 and S2 cells. Transfected cells

were harvested with enzyme-free cell dissociation buffer (Gibco), centrifuged, and suspended in assay

media (DMEM with 0.1% BSA for mammalian cells, unsupplemented SFX-100 insect medium for S2). The

aequorin cofactor coelenterazine-F (Nanolight) was added (5 µM final concentration) to the cell

suspension, in the dark, with gentle stirring for 2 to 3 hours before assay. For water-insoluble compounds

like DEET an emulsion was prepared by adding a percentage of DEET indicated, vol/vol to assay media,

vortexed, and bath-sonicated for at least 15 min, but up to one hour. Cells (100 µl, at approx. 50,000 cells)

were injected into a 96 well plate containing 100 DEET µl emulsion, and resulting luminescence recorded

on a BMG Labtech Lumistar Omega plate reading luminometer. Thapsigargin was added to the cell

suspension were indicated to a final concentration of 1 µM, 10 minutes before assay.

Chemical Informatics

Machine learning predictions of insect repellency

Physicochemical features for figures 1-2 were selected using Sequential Forward Selection (SFS) (Haddad

et al., 2008) from 3,224 using Dragon software (Talete). This algorithm is an iterative search over the

chemical feature pool that seeks to maximize the correlation between chemical feature values and, here,

the % repellency. These chemical features were subsequently used to fit a support vector machine (SVM)

with a radial basis function kernel (RBF). The implementation of the support vector machine algorithm is

from the e1071 R package. We performed 20 independent 5-fold cross-validations followed by a receiver

operating characteristics (ROC) analysis to analyze the performance of the repellency prediction method.

The trained SVM then ranked both a library obtained from eMolecules (http://www.emolecules.com)

(>400,000) and natural odor libraries (Supplementary Material) (>3,000) by the repellency score estimates

and the set of optimally predictive chemical features. The EPI Suite

(http://www.epa.gov/oppt/exposure/pubs/episuite.htm) was used to calculate predicted LogP and Vapor

Pressure values.

For Figures 3, 5, S1-S2, we updated the methods and algorithms to further optimize the pipeline.

Physicochemical features were selected using AlvaDesc software, which offers > 5000 chemical features.

Optimal chemical features were then selected using cross-validation and recursive feature elimination

algorithm (CV-RFE). The RFE CV algorithm involves iterative partitioning of the data and the selection of

feature subsets (n_1, n_2, …, n_{i+1}; where "n" is the size of the feature subset and "i" is an arbitrary indexer).

Given a data partition and n features, an arbitrary ML algorithm is fit, importance is assigned to the feature

set and the model of n features predicts a test partition, that is, data excluded from training,

differentiating the CV-RFE from RFE alone. In this study, the ML algorithm was a support vector machine

(SVM) or a random forest. Over 100s of iterations the algorithm converges on the optimal (n) number of

features to maximize predictive success. Additionally, for every round the contribution of each chemical

feature is assessed, giving rise to an aggregate rank; namely, the frequency at which a feature has been

identified as important to the predictive success. Again, distinguishing the standard RFE algorithm from

CV-based adaptation. As for computing importance, the random forest algorithm internally tests the

contribution of the features. It accomplished this by fitting multiple decision trees on bootstrap samples

of the data. Consequently, a percentage of data has been excluded and serves as a test set. By predicting

the test set with and without permuting or shuffling the values of a given feature, the algorithm can then

assign an importance according to the % increase in error that is observed after permuting the feature.

The support vector machine (SVM) algorithm does not natively perform a comparable

bootstrapping procedure as random forest and therefore does not generate a test set to assess feature

importance. In this case, an external metric must be supplied to assign the importance. Depending on the

type of fit, classification or regression, this is a pseudo R^2, based on non-linear regression, or the AUC for

the feature and the outcome that is being predicted (e.g., % repellency). For additional details, refer to

the documentation provided with caret package in R.

Machine learning predictions of Ca^{2+} mobilization

Due to the smaller number of observations (datapoints or chemicals) we opted for an ensemble

method that could take advantage of more of the available data. Random forest fits multiple decision

trees on bootstrap samples of the data, thereby reducing the tendency to overfit to one or few chemicals

in the dataset. As compared to the SVM, it has two simple parameters (mtry and ntrees) and the tuning

grid (or number combinations of these parameters that need to be tested to optimize performance tends

to be small). Accordingly, it is an ideal approach when little is known about the activity being predicted.

To further reduce overfitting, we identified a small set of physicochemical features on different

subsets of the data and then randomly sampled from this pool. By doing so, the features are slightly

underfit, which is the better approach, again, when less is known about the activity being predicted.

Additionally, we used two different methods and metrics to independently verify that

physicochemical features were indeed predictive of the activity. First, a regression-based approached

predicting the raw assay activities. Second, a classification-based approach where the top ~40% of the

activity distribution was converted into an "Active" label, assessing predictive success by ROC analysis.

Finally, we fit classification models that were otherwise identical but trained on shuffled labels. This

ensured that the chance level-performances significantly differed.

Machine learning predictions of odor perceptual qualities

We used methods that were developed previously and extensively validated for prediction of human odor

characters from chemical structures and are described there (Kowalewski et al., 2021; Kowalewski & Ray,

2020b). Briefly, perceptual training data was from a study published in a reference book: ATLAS of Odor

Character Profiles (Dravnieks, 1985). In the study, panelists supplied ratings for ~150 odorants and

mixtures across 146 possible descriptors; the ratings are reported as the % usage. The % usage refers to

the portion of panelists that rated an odorant (1-5) using a particular descriptor (1-146), indicating its

relevance to the odorant. It is on the scale 0-100, with the maximum of 100% suggesting every participant

found the descriptor to be relevant. The top perceptual descriptors were selected by thresholding the

predictions based on the mean value in the training data. For each descriptor in the training data,

thresholds were computed as 2x the mean. These thresholds were then applied to the predicted

repellents to generate a set of exemplar descriptors. Subsequently, these exemplars were converted into

a descriptor frequency table, where the frequency is the total number of occurrences of a descriptor. The

data and word cloud plot were generated using the R programming language, with additional functionality

provided by the Wordcloud package in R.

RESOURCE AVAILABILITY

Lead Contact

Anandasankar Ray, anand.ray@ucr.edu

Materials Availability

No new materials were generated in this study.

Data and Code Availability

Data used in the analyses is publicly available from the references cited, and the data generated in this

manuscript are supplied within the figures. Any additional data associated with this manuscript is

also available in other formats on request from the corresponding author and lead contact.

The code developed for this study is available from the corresponding author for non-commercial use

upon request. Use of the code may be subject to a standard Material Transfer Agreement (MTA) or

specialized licensing agreement through University of California Riverside's Technology Transfer Office,

as patent applications related to this software are currently pending.

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AUTHOR CONTRIBUTIONS

454   J.K., S.M.B., R.A. and A.R. conceived the study. J.K. and S.M.B. planned and executed the computational

455   analysis. R.A. planned and executed the cell line experiments and J.A. performed the insect behavior

456   experiments. J.K. and A.R. wrote the manuscript with contributions from others.

DECLARATION OF INTERESTS

J.K., S.M.B., and A.R. are listed as inventors on various patent applications filed by the university. A.R. is

the founder of Sensorygen Inc., that has licensed some of the intellectual property, and A.R. is also the

founder of startup companies: Remote Epigenetics, and Sensorygen. Sensorygen has licenses to some of

the IP from UCR and is involved in finding new flavors and insect repellents.

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 FIGURE LEGENDS

Figure 1. A chemical informatics pipeline to screen for pleasant smelling insect repellents

A, Overview of the cheminformatics pipeline used to identify novel DEET-like ligands from a larger

chemical space. Summary table of top 10 physicochemical features that optimally corelate with

repellency. Receiver-operating-characteristic curve (ROC), which indicates the success in predicting test

repellents. The performance is the average over 100 80/20 splits, where 80% of the data is for training

and 20% for testing. The mean area under the curve (AUC) is provided. Representative structures from

the top 150 predicted repellent compounds.

Figure 2. Models successfully predict odor perceptual qualities of repellents in humans

A, Validation of models predicting the odor perceptual qualities (146 perceptual descriptors) of a set of

compounds that were screened for repellency in Figure 3A. The performance metric for the validation is

the area under the ROC curve, which assesses the rate of correctly predicted perceptual descriptors

relative the rate of incorrectly predicted descriptors. Dashed line is the average Area Under the Curve

(AUC). **B,** Sample ROC curves for select compounds as well as the ROC curve across all compounds (e.g.,

shown as separate curves in the ROC plot as well as those not shown but whose identities appear in Figure

2A); the respective AUC values are reported inside the plot. **C,** The top 25 perceptual descriptors for the

predicted repellents that underwent experimental validation are displayed as a word cloud. The font size

of the descriptor is scaled relative to frequency.

Figure 3. Experimental validation of predicted insect repellents show a high success rate several of

which are pleasant smelling

A, Representative still photograph of landing assay of female *Aedes aegypti* mosquitoes on filter paper

covered heat block over 5 mins (overlay of all frames), and **B,** mean Repellency Coefficient for each of the

indicated compounds (by CAS numbers) tested at ~0.2mg/cm^2, N=2-4 trials for each, ~20 mosquitoes/trial.

For DEET, N=23 trials. CAS numbers ending in a ".1" are compounds tested at a second lower amount of

~0.06mg/cm^2.

C, Mean Preference index of *Drosophila* adults to repellents at three different concentrations dissolved in

acetone in a Two-choice trap assay baited with 10% Apple Cider Vinegar measured after 48 hrs. N = 7-10

trials each treatment at 48 hrs, 10 flies/trial, error bars = s.e.m.

Figure 4. Some, but not all insect repellents increase Ca^{2+} mobilization in a human cell line

DEET induces a non-specific calcium response *in vitro*. **A.** Dose response curves for three mammalian cell

lines (HEK293. CHOK1, NG108) and one insect cell line (S2). EC50 and Hill coefficients for each cell line are

given below. **B.** Representative DEET induced calcium kinetics in HEK293 cells. **C.** HEK293 cells treated

with thapsigargin 10 min before assay to deplete intracellular calcium stores no longer respond to DEET,

suggesting DEET response is dependent on intracellular calcium. Error bars are standard deviation. **D. F,**

Luminescence of the Ca^{2+} indicator at two different concentrations for predicted and previously known

repellent chemicals, scaled relative to DEET.

Figure 5. Screen of 10+ million chemicals predicts several insect repellents with different smells

A, Tabulated predicted % repellency (relative to DEET) from a library of 10+ million chemicals, filtered to

the top values. The best matching known repellent is displayed alongside the Euclidean distance, the

predicted LD50 and Ca^{2+} mobilization. **B**, heatmap organizing top predictions according to estimated

perceptual qualities.

Supplemental Figure S1

A, Overview of the approach to generate and validate the machine learning models. Here, a model-

averaged prediction is made, where each model has access to different physicochemical features and is

trained on different combinations of training set chemicals. **B,** The R^2 values over the 100 train/test splits,

averaged into 10 bins. **C,** The average classification success is reported over 100 train/test splits, assessed

by the area under the ROC curve. Active labels (positive cases) were assigned as chemical scoring in the

top ~40% of Ca^{2+} values. This is compared to shuffling the labels before training, reported as "Shuffle."

Descriptor Symbol	Descriptor Definition
B05[C-N]	Presence/absence of C-N at topological distance 5
piID	conventional bond order ID number
N-072	RCO-N< />N-X=X
nRCONR2	number of tertiary amides (aliphatic)
RDF035v	Radial Distribution Function - 035 / weighted by van der Waals volume
EEig08r	Eigenvalue 08 from edge adj. matrix weighted by resonance integrals
D-Dr06	distance/detour ring index of order 6
Mor10p	3D-MoRSE - signal 10 / weighted by atomic polarizables
BAC	Balaban centric index
EEig14d	Eigenvalue 14 from edge adj. matrix weighted by dipole movements

Figure 1

Figure 2

Figure 3

Figure 4

A

Image	ZINC_ID	Score	LD$_{50}$ mg/kg	Ca^{2+} mobilization	Image	ZINC_ID	Score	LD$_{50}$ mg/kg	Ca^{2+} mobilization
	ZINC000001394030	110.9316	1293.0062	0.9054749		ZINC000008609700	110.2547	1287.3842	0.5077889
	ZINC000034195873	110.5746	711.4268	0.7932151		ZINC000253417706	109.7302	933.5598	0.8336481
	ZINC000005327991	110.4276	883.9981	0.7673523		ZINC000034195871	109.6462	684.8090	0.9535496
	ZINC000094992683	110.4188	1067.0473	0.9159867		ZINC000000506855	109.5113	1299.5812	0.7571195
	ZINC000001735256	110.4169	883.8132	0.7749915		ZINC000191321911	109.5082	989.3195	1.1121364

B

Figure 5