

Applying IBM Watson cognitive computing to identify drugs with potential for treating L-DOPA-induced dyskinesia

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Background

Current treatment options for L-DOPA-induced dyskinesia (LID) are limited. Repurposing compounds with regulatory approval is an attractive approach to accelerate availability of novel therapeutic options. IBM Watson for Drug Discovery (WDD) is a cognitive computing platform that extracts domain specific text features from vast quantities of published literature to identify existing and infer novel connections between entities of interest, including drugs.

Objectives

To use WDD to analyze published abstracts of a set of compounds known to prevent LID and apply machine learning to rank a candidate set of compounds according to similarity of linguistic context and thereby discover compounds with potential for treating LID.

Methods

15 compounds, of a range of classes, with demonstrated ability to reduce LID pre-clinically or in clinical trials were identified as known compounds.

| Class of compound | Compound name |
|----------------------------------------------|---------------|
| NMDA receptor antagonists | |
| Cannabinoid receptor 1/2 partial agonist | |
| mGlu5 NAMs | |
| AMPA antagonists | |
| alpha2 adrenergic antagonists | |
| 5-HT1A agonists | |
| 5-HT1B agonists | |
| alpha 7 containing nicotinic agonists | |
| alpha 4 beta 2 containing nicotinic agonists | |
| mu/k-opioid receptor antagonist | |

Table 1. Known compounds

Candidate compounds were filtered from the entire DrugBank database of 8261 entries including 2021 FDA-approved small molecule drugs, 233 FDA-approved protein/peptide drugs, 94 nutraceuticals and over 6000 experimental compounds (<https://www.drugbank.ca/>). Dopamine agonists and compounds with <5 published abstracts were removed, leaving 3539 final candidates. WDD analyzed ~1.3 million Medline abstracts and then, using machine learning, created a predictive model to rank candidate compounds based on semantic similarity to the known set (Figure 1).

Figure 1. Predictive analytics methodology

Each compound is converted to a vector, or "semantic fingerprint," representing the weighted frequency of the words and phrases in all abstracts that mention it. From these vectors, a distance matrix relating every compound in the known and candidate lists is generated. A semi-supervised learning approach, graph diffusion (1), is then used to rank candidate compounds by similarity to the overall 15 known compounds.

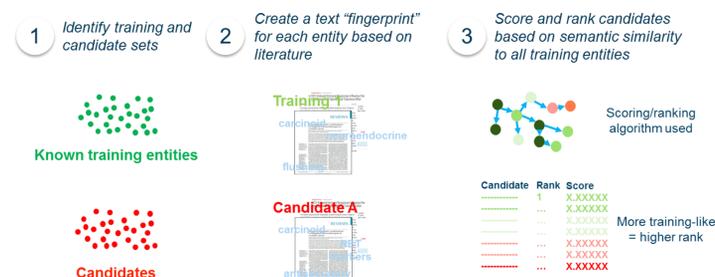


Figure 2. Leave-one-out cross-validation

The analysis is run 15 times, each time with one entity from the known set excluded and added to the candidate set. Wilcoxon p-value (4.5×10^{-11}) and area under the Receiver-Operator Characteristics curve (AUC=0.72) demonstrate that the semantic models Watson has created from the known drugs have predictive power over each other compared to the candidate set.

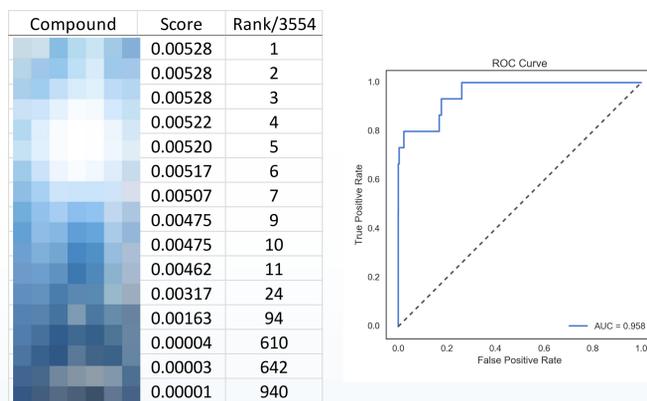


Table 2. Ranking of known compounds

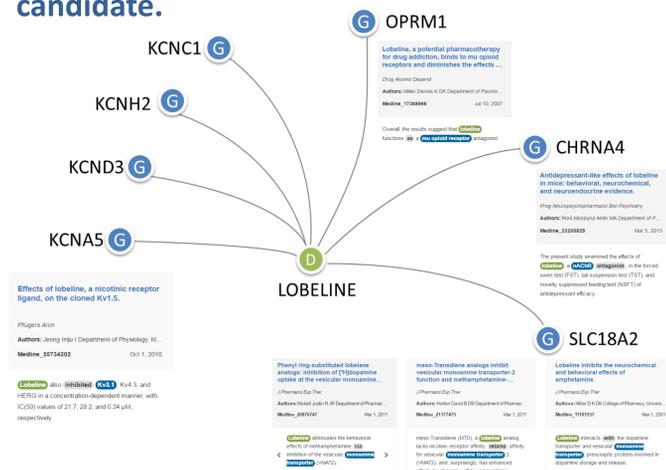
Retrospective analysis

To further validate WDD's methodology, the analysis was first restricted to abstracts published up to and including 2013. 3 known drugs with abstracts first demonstrating their antidyskinetic effects published after 2013 were omitted from the known set. Using only 7 and 4 available abstracts respectively, WDD ranked 2 of these highly, 111th, and 194th (top 5.5%) out of 3527 drugs. The third, ranked 857th (top 25%).

Prospective Analysis: Results

The top 50 ranked candidates were explored in WDD to determine their relationship with each drug in the known set, and reveal the biological connections that may contribute to their predicted antidyskinetic actions.

Figure 3. Case study of a highly ranked candidate.



| Relationship type | Verb | Target (Canonical) | Target (Text) | Target type | Sentence fragment |
|--------------------------------|-------------|---------------------------------------------------|---------------------------------------------------|-------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Prepositional | as | OPRM1 | mu opioid receptor | GENE | Overall, the results suggest that lobe-line functions as a mu opioid receptor antagonist. |
| RegulationNegative | antagonist | CHRNA4 | nAChR | GENE | The present study examined the effects of lobe-line, a nAChR antagonist, in the forced swim test (FST), tail suspension test (TST), and novelty-suppressed feeding test (NSF) and found that lobe-line significantly reduced the behavioral effects of methamphetamine via inhibition of the vesicular monoamine transporter (VMAT2). |
| Prepositional | with | SLC18A2 | monoamine transporter | GENE | Lobe-line interacts with the dopamine transporter and vesicular monoamine transporter, presynaptic proteins involved in dopamine release. Lobe-line attenuates the behavioral effects of methamphetamine via inhibition of the vesicular monoamine transporter (VMAT2). |
| RegulationNegative/Association | inhibited | KCNH2 | HERG, KCNC1, Kv3.1, Kv4.3, Kv1.5 | GENE | Lobe-line also inhibited Kv3.1, Kv4.3, and HERG in a concentration-dependent manner. |
| RegulationNegative | attenuated | nicotine-induced hyperactivity | nicotine-induced hyperactivity | NA | Lobe-line attenuated nicotine-induced hyperactivity when both drugs were administered respectively. |
| RegulationNegative | inhibit | blood-brain (BBB) transporter | the blood-brain barrier BBB amine transporter | NA | Lobe-line is a potential smoking cessation drug with affinity for the alpha2B nicotinic acetylcholine receptor and may inhibit the blood-brain barrier (BBB) amine transporter. |
| Regulation | mediate | lobe-line | the agonist effects of lobe-line | NA | Although the alpha2beta2 receptor is unlikely to mediate the agonist effects of lobe-line, our results indicate that lobe-line does interact with the nicotinic receptor in a novel fashion. |
| Association | desensitize | (inhibition) | receptors when applied alone inhibition | NA | Thus, lobe-line can apparently desensitize receptors when applied alone (inhibition) whereas its binding to a second agonist site with the first one already occupied by acetylcholine leads to channel opening (potentiation). |
| RegulationNegative | reduces | epileptic seizures | epileptic seizures | NA | Hence, we may propose that lobe-line reduces epileptic seizures by enhancing the GABA release supporting the GABAergic mechanism. |
| RegulationNegative | reduced | lobe-line | the effects of lobe-line | NA | The depressant effects of lobe-line on the visual response could not be reversed by the GABA(A) antagonist bicuculline, but the GABA(B) antagonist CGP 35348 reduced the effects of lobe-line. |
| RegulationNegative | decreased | locomotor activity and body temperature in mice | locomotor activity and body temperature in mice | NA | Lobe-line, at the time of maximal effect, dose-dependently produced motor impairment and decreased locomotor activity and body temperature in mice after s.c. |
| Association | produced | hypoactivity | hypoactivity | NA | Lobe-line produced hypoactivity in total horizontal activity and center distance travelled. |
| RegulationPositive | induced | stereotyped behaviors | any stereotyped behaviors | NA | In saline challenge groups, the doses of lobe-line examined did not affect spontaneous locomotion nor induced any stereotyped behaviors. |
| RegulationNegative | attenuates | the behavioral effects of methamphetamine | the behavioral effects of methamphetamine | NA | Lobe-line attenuates the behavioral effects of methamphetamine via inhibition of the vesicular monoamine transporter (VMAT2). |
| RegulationPositive | augments | cocaine-induced hyperactivity | cocaine-induced hyperactivity in rats | NA | Lobe-line augments and inhibits cocaine-induced hyperactivity in rats. |
| Regulation | alters | presynaptic dopamine storage and release in brain | presynaptic dopamine storage and release in brain | NA | Lobe-line has high affinity for nicotinic receptors and alters presynaptic dopamine storage and release in brain. |
| Regulation | alters | dopamine | dopamine function | NA | Reevaluation of the mechanism by which lobe-line alters dopamine function reveals that its primary mechanism is inhibition of dopamine uptake and promotion of dopamine release from the storage vesicles within the presynaptic terminal, via an interaction with the tetraabenazine-binding site on the vesicular monoamine transporter (VMAT2). |
| RegulationNegative | inhibit | DA uptake synaptic vesicles | DA uptake into synaptic vesicles | NA | These results suggest that lobe-line specifically interacts with DTBZ sites on VMAT2 to inhibit DA uptake into synaptic vesicles. |
| Localization | displaced | 3H]cytisine cortical | 3H]cytisine binding to rat cortical membranes | NA | Lobe-line displaced 3H]cytisine binding to rat cortical membranes with a mean inhibition constant (Ki) value of 16.0 nM, while the lobe-line analogs CRM-1-13-1 and CRM-1-32-1 exhibited values of 15.0 and 5.1 microM, respectively. |
| Regulation | mediated | lobe-line | the effect of lobe-line | NA | While DMPP-induced 5-HT release can be linked to a non-classical nAChR activation ([Ca2+]i-dependence), the effect of lobe-line was likely mediated by uptake carriers. |
| RegulationNegative | attenuates | behavior deficits in animals | behavior deficits in animals | NA | Lobe-line shows protective effects against MPTP-induced dopaminergic neuron death and attenuates behavior deficits in animals. |

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Screen shots from WDD illustrating a selection of the gene relationship network for Lobe-line (ranked 39/3539) and a table of a selection of the unconstrained relationship network.

Discussion

We have demonstrated the feasibility of using WDD to rank compounds based on their predicted potential to reduce LID. Further studies will be required to validate these predictions in animal models of LID. As well as identifying compounds with known and plausible antidyskinetic mechanisms of action, future work will address the antidyskinetic potential of compounds identified by WDD with a novel/unknown mechanism of action and test the power of WDD to uncover novel strategies to tackle this debilitating condition.

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