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6	Deep Q-learning for the selection of optimal isocratic
7	scouting runs in liquid chromatography
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30 ABSTRACT

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32 An important challenge in chromatography is the development of adequate separation methods. Accurate retention models can significantly simplify and expedite the development of adequate 33 separation methods for complex mixtures. The purpose of this study was to introduce 34 reinforcement learning to chromatographic method development, by training a double deep O-35 learning algorithm to select optimal isocratic scouting runs to generate accurate retention 36 models. These scouting runs were fit to the Neue-Kuss retention model, which was then used 37 38 to predict retention factors both under isocratic and gradient conditions. The quality of these 39 predictions was compared to experimental data points, by computing a mean relative 40 percentage error (MRPE) between the predicted and actual retention factors. By providing the reinforcement learning algorithm with a reward whenever the scouting runs led to accurate 41 42 retention models and a penalty when the analysis time of a selected scouting run was too high (> 1h); it was hypothesized that the reinforcement learning algorithm should by time learn to 43 44 select good scouting runs for compounds displaying a variety of characteristics. The reinforcement learning algorithm developed in this work was first trained on simulated data, 45 46 and then evaluated on experimental data for 57 small molecules - each run at 10 different fractions of organic modifier (0.05 to 0.90) and four different linear gradients. The results 47 48 showed that the MRPE of these retention models (3.77% for isocratic runs and 1.93% for 49 gradient runs), mostly obtained via 3 isocratic scouting runs for each compound, were comparable in performance to retention models obtained by fitting the Neue-Kuss model to all 50 (10) available isocratic datapoints (3.26% for isocratic runs and 4.97% for gradient runs) and 51 52 retention models obtained via a "chromatographer's selection" of three scouting runs (3.86% 53 for isocratic runs and 6.66% for gradient runs). It was therefore concluded that the reinforcement learning algorithm learned to select optimal scouting runs for retention 54 55 modeling, by selecting 3 (out of 10) isocratic scouting runs per compound, that were 56 informative enough to successfully capture the retention behavior of each compound.

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- 58

59 Keywords: method development; retention models; machine learning; reinforcement

60 learning; deep q-learning;

61 1 INTRODUCTION

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63 One of the main challenges in (liquid) chromatography is to obtain adequate methods to 64 separate complex samples. Method development is still often done based on a trial-and-error 65 approach, leading to lengthy optimization procedures. Decision-algorithms can be used to help 66 find adequate separations for complex samples (1-8). These algorithms need to be able to 67 predict future runs and select the correct runs to produce the most desirable outcome, which 68 means accurate retention models need to be incorporated into the algorithms. Retention models 69 can generally be divided into two types: quantitative structure retention relationship (QSRR) 70 models, that model retention times of compounds for specific separation conditions based on 71 their molecular descriptors (9-16) and empirical models, that model retention times of compounds based on experimental instrument or system variables, e.g. the fraction of strong 72 73 eluent in the mobile phase (17-24). The two types of retention models are complementary and 74 usually answer different questions. For instance, QSRR models are commonly developed to 75 improve the identification of known analytes, while empirical retention models are rather 76 developed to improve the separation of complex mixtures of known or unknown compounds. 77 As the present study focuses on improving separation methods for compounds for which 78 molecular structures are not always available, empirical retention models will be used.

79

81

80 An often employed empirical retention model is the Neue-Kuss model (21, 26-28):

$$k(\varphi) = k_w (1 + S_2 \varphi)^2 \exp\left(-\frac{S_1 \varphi}{1 + S_2 \varphi}\right)$$

82 83

84 Wherein φ is the fraction of strong eluent in the mobile phase, $k(\varphi)$ the retention factor for a 85 specific ϕ and k_w, S₁ and S₂ are retention parameters that need to be fit to experimental retention factors to yield a retention model for a given compound. More specifically, kw refers to the 86 87 retention factor of a compound for $\phi=0$ (in reversed-phase liquid chromatography (RPLC) this 88 corresponds to a purely aqueous mobile phase), S_1 refers to the slope and S_2 to the curvature of 89 the correlation between k and φ . Since the Neue-Kuss model requires three parameters for the prediction of the retention factor (k_w , S_1 and S_2), at least three datapoints (*i.e.*, experimental 90 91 retention factors obtained at three different mobile phase compositions) are needed as input. In 92 RPLC, the polarity of a compound is one of the main factors influencing its retention behaviour. 93 Since compounds can have large differences in polarity, the selection of the scouting runs, *i.e.*, 94 the selection of the mobile phase compositions that need to be selected to obtain adequate 95 retention models, is an important problem as the most informative conditions can strongly 96 differ for each compound. This importance is emphasized making the economic consideration 97 that the number of scouting runs should preferably be kept as low as possible. Thus, there is a 98 clear need to develop algorithms that can select optimal scouting runs leading to accurate 99 retention models.

100

101 Decision algorithms for chromatographic applications, such as method development, have been 102 studied since the 1980's. These algorithms, also sometimes referred to as *expert systems*, 103 focused on different aspects of the chromatographic workflow: method selection (28,29), 104 mobile phase selection (30,31), mobile phase/selectivity optimization (32,33), system 105 optimization (34,35) and retention optimization (36-39). All these algorithms aimed at 106 improving the decision-making process for method development and reducing the extent of 107 trial-and-error that is commonly practiced. However, in the late 90's these expert systems lost 108 their popularity due to the fact that the high expectations for these systems were not met, and 109 the computational power available at that time was inadequate.

(1)

110 Traditional approaches, like those used in the 80's and 90's, usually take the form of predefined 111 rules and if-else statements, which are directly programmed/handcrafted by the software engineer and experts in the field. The inherent problem of this approach is its great difficulty 112 113 to generalize to new data or scale to large, diverse datasets. Algorithms like machine learning 114 (40) can, to some extent, solve the problem of generalization by letting the machine program 115 itself (learn its own rules) via highly parameterized and regularized function approximators 116 that are trained on a so-called training dataset. Supervised machine learning algorithms have 117 been around for many decades (41-44) but came to proliferate in the late 2000's when the 118 availability of data in both private and public repositories grew considerably and sufficiently 119 powerful computer hardware became readily available. This family of algorithms only needs 120 to be supplied with input and output pairs, and everything in-between (mapping from input to 121 output) is learned by the algorithm itself. Given an input, supervised learning algorithms learn 122 how to minimize the error between their prediction and the target (the associated output for 123 that input). A typical application of supervised learning is classification, in which, given an input, the algorithm outputs the correct class. However, the limitation of supervised learning is 124 125 the need to label every input, which is often either very time-consuming or impossible to 126 actualize. Unsupervised learning deals with situations where no labels are required, with a 127 typical application being clustering. Instead of directly getting a prediction from some input, it 128 clusters the input data and finds patterns in it. Either of these two paradigms is important, but 129 neither fulfills the criteria for intelligent decision making as required in method development. 130 In method development, especially when dealing with novel or unknown compounds, a 131 prediction of the next step to perform is needed, but the effects of the steps are unknown and 132 impossible to plan.

133

134 Reinforcement learning is a machine learning paradigm that deals with sequential decision 135 making (45,46). In contrast to supervised learning, where each prediction is *independent* of the 136 others, decisions are sequential and *dependent* on each other. This aligns well with method 137 development, which is clearly a sequential procedure (cfr. the trial-and-error approach when 138 method development is carried out in its most simple form). Furthermore, in reinforcement 139 learning, there is no need for input-output pairs as in supervised learning, because no label is 140 needed to teach the algorithm to perform well on a given task (which is of great importance as 141 labels are commonly unavailable in method development). Instead of a label, a scalar reward 142 signal reinforces a desirable behavior; *i.e.* the reinforcement learning algorithm learns a 143 mapping from some input to some desirable output by receiving feedback based on the output. 144 The goal of the reinforcement learning algorithm is not to optimize itself towards accurate label 145 predictions, but instead maximize the cumulative future reward (*i.e.* learn to optimally perform 146 a specific task). Due to its generality, reinforcement learning algorithms have been widely 147 utilized and studied in many applications and areas of research and industry, including robotics 148 (44–49). Besides its many applications, reinforcement learning is possibly most known for its 149 achievements in attaining superhuman level performance in board games, such as Chess and 150 Go (53–55), and in video and computer games such as Atari (56), Starcraft II (57,58) and Dota 151 2 (59).

152

In the present study, the possibilities of reinforcement learning are investigated for a basic chromatographic problem: selecting optimal isocratic scouting runs in RPLC to fit retention models. Specifically, the aim is to train a reinforcement learning algorithm to select a minimum number of scouting runs that allow to obtain accurate retention models for a diverse set of compounds. The present study is considered a simple first step in introducing reinforcement learning to the area of chromatography.

160 **2** THEORY

161

162 Reinforcement Learning is a goal directed paradigm in which a computer program, called the 163 agent, is continuously interacting with the environment. The environment is the world in which 164 the agent resides and encapsulates the task the agent tries to solve. By performing actions, the 165 agent receives a numerical feedback from the environment based on the quality of the chosen 166 action. Over time, the agent learns to optimize the actions it takes by trying to maximize the 167 future rewards. Typically, reinforcement learning uses a Markov Decision Process (MDP) as 168 its mathematical model. An MDP is a 4-tuple $\langle S, A, T, R \rangle$, where S is the state space, A the 169 action space, $T(s, a, s') = \Pr(s' \mid s, a)$ the probability that action a in state s will lead to state s', 170 and R(s, a, s') the immediate reward (r) received when action (a) taken in state (s) results in a 171 transition to state (s'). The problem of solving an MDP is to find a policy (π) (i.e., mapping 172 from states to actions) that maximizes the accumulated reward (i.e., the sum of all scalar reward 173 signals). A typical interaction goes as follows: given a state s, the agent takes an action a, which 174 causes the environment to change its state to a new state s'. In addition to the new state s', the 175 environment provides a specific reward signal, indicating how good it is in the new state. 176 Moving across states by performing an action is known as a *transition*, and a sequence of 177 transitions is called a *trajectory*. Specifically, in this study, a trajectory (also referred to as an 178 episode) is defined as a sequence of transitions and ends with a *termination* signal indicating 179 that no more scouting runs are to be run for a given compound.

180 The state space and action space of the environment define the space in which the 181 reinforcement learning algorithm (or the agent) can perform intelligently. The state of the 182 environment can be represented in many ways, everything from video/image input (a 183 continuous state space), to a vector containing the positions of marks on a tic-tac-toe board (a 184 discrete state space). Similarly, the action space may also be continuous and/or discrete. The 185 environment also sets restrictions for what actions are legal – for instance, is the agent allowed 186 to adjust the percentage of modifier to any percentage? Or is it only allowed to change it 187 between certain discrete, predefined, ranges?

188 To formalize the learning procedure, the agent tries to learn a policy – what action to take given 189 a certain state. The ultimate goal of the agent is to learn a policy π that maximizes the expected 190 cumulative future reward (known as the expected return) over a trajectory τ . This optimal 191 policy denoted as π^* is defined as:

192 193

 $\pi^* = \arg \max_{\pi} E_{\tau \sim \pi}[R(\tau)], \qquad (2)$

194

195 where
$$R(\tau) = \sum_{t=0}^{\infty} \gamma^t r_t$$
 (3)
196

197 $E_{\tau \sim \pi}[R(\tau)]$ denotes the expected return of trajectory τ following policy π and $\arg \max_{\pi}$ 198 specifies the policy maximizing the expected return. $\gamma \in (0,1)$ is known as the *discount factor* 199 that discounts the reward r depending on how far off in the future the reward was obtained. If 200 the discount factor is smaller than one, it is more valuable to get rewards early on rather than 201 later.

202

For the agent to judge the quality of a given state in an environment, it learns its value through a value function. The value function denotes the desirability of being in a state at a given point in time. In the present study, the *optimal action-value function* Q^* will be used – which expresses the expected return if the agent starts in a state s, takes an action a, and then acts according to the optimal policy forever after:

208
$$Q^*(s,a) = \max_{\pi} E_{\tau \sim \pi}[R(\tau)|s,a]$$
(4)

210 \max_{π} indicates the policy yielding the maximum expected return and $\tau \sim \pi$ indicates that the trajectory is sampled according to the policy. Importantly, Q^* can be rewritten as a recursive 211 212 function, called the Bellman equation for Q^* (60):

214
$$Q^*(s,a) = E_{s' \sim P} \left[r(s,a,s') + \gamma \max_{a'} Q^*(s',a') \right]$$
(5)

215

223

225

213

 $s' \sim P$ indicates that the next state is sampled according to the environment's transition rules, 216 r(s, a, s') is the reward received in the next state s' by taking action a in state s, and 217 $\gamma \max_{a'} Q^*(s', a')$ is the discounted maximum expected return in the next state s'. The 218 expected return Q^* , also referred to as the Q-value, of each state-action pair can be computed 219 - allowing for the optimal action a^* in a given state to be obtained via: 220 221

222
$$a^*(s) = \arg \max_a Q^*(s, a)$$
 (6)

224 arg max_a specifies the action which maximizes $Q^*(s, a)$ and yields the highest Q-value.

226 The optimal action-value function (Q^*) is not known at the start and needs to be solved. A 227 common approach to solve Q^* is via *Q*-learning (61). The *Q*-learning algorithm incorporates 228 the Bellman equation for Q^* to iteratively update Q: 229

230
$$Q_{i+1}(s,a) = E_{s'} \left[r + \gamma \max_{a'} Q_i(s',a') | s, a \right]$$
231 where $Q_i \to Q^*$ when $i \to \infty$
(7)

232

Υi

233 Q is known as the Q-function, which approaches or represents Q^* at any given time. i indicates 234 iteration *i*, or transition *i*. By experiencing states and actions, O can iteratively be updated until 235 it reaches Q^{*} or is close enough to it. The agent will select actions based on an action selection 236 mechanism. A typical method for action selection is ϵ -greedy and is defined as: 237

238

$$a \leftarrow \begin{cases} \arg \max_a Q(s, a) & \text{with probability } 1 - \epsilon \\ \text{random action} & \text{with probability } \epsilon \end{cases}$$

239

240 In essence, the agent will choose to perform a random action with probability ϵ , and will select 241 the optimal action (based on what the agent knows about the environment so far) with a 242 probability 1- ϵ . Through this continual interaction, the agent will eventually learn which 243 actions to perform at each state.

244

245 However, a problem with this tabular approach is that it is impractical, because the state space 246 and action space in most interesting problems are often too large. This approach may not 247 converge to Q^* and lacks generalization capabilities (only able to yield accurate expected 248 returns given actions and states that it has experienced previously). In reinforcement learning, 249 generalization is typically solved by some form of function approximation by representing the 250 value function as a linear function or as a neural network. A major breakthrough in 251 reinforcement learning was the use of the representational power of deep neural networks 252 instead of tabular value functions which led to the creation of Deep Q Networks (DQN) (56) 253 and spurred an entire research field called Deep Reinforcement Learning. 254

Specifically, a function approximator, in this case a deep neural network, approximates $Q^*: Q_\theta \approx Q^*$. The parameterized action-value function Q_θ is thus a neural network, referred to as the *Q*-network, which receives as input a state *s* and outputs the expected return for each possible action in that state. The parameters θ of the *Q*-network are estimated by *minimizing* an *objective function* [with respect to θ], which aims to yield an accurate approximation of Q^* : 260

- 261 $\min_{\theta_i} \left[\left(r(s_i, a_i, s_i') + \gamma \max_{a_i'} Q(s_i', a_i', \theta_i^-) \right) Q(s_i, a_i, \theta_i) \right]^2$ (8)
- 262

263 \min_{θ_t} indicates that θ is being adjusted at each iteration (*i*) to minimize the objective function. 264 The objective function is the mean squared error between the target (left term between brackets) 265 and the prediction (right term between brackets). For stability purposes, the *target* network has 266 its parameters θ^- held fixed when minimizing the objective function to not have a continually 267 moving target. The method is referred to as *DQN*, and the minimization procedure referred to 268 as *training* – which is an iterative procedure that iteratively tries to minimize the objective 269 function over the course of thousands of episodes. 270

271 To better understand the concept of reinforcement learning from a higher level, and to tie it to 272 the process of method development, let us consider the agent-environment interaction. This 273 can be thought of as an analyst performing liquid chromatography (LC). If the environment is 274 the LC with its internal physics and chemistry (for example the interactions of analytes with 275 the stationary phase inside the column), then the agent would be the analyst, interacting with 276 the LC. The analyst merely sees a representation of the interaction between a mixture of 277 compounds and the mobile/stationary phase. This representation will take the form of a 278 chromatogram (or retention times), which is the output of the LC. Based on this output, the 279 analyst decides the next action, and as a consequence of this new action, a new representation 280 of a state (new output) is presented to the analyst, and so on. These abstractions are important 281 to understand the roles of the agent and the environment and how they interact, as well as the 282 important differences between them.

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285 3 MATERIALS AND METHODS

286

287 **3.1** Chemicals

288

289 The compounds used in this study, together with their structure, Log P and pKa values, stock 290 solvent and supplier, are mentioned in Table S-1 in the Supporting Information. Structures 291 were drawn in Marvin Sketch (v20.9.0, 2020, ChemAxon (www.chemaxon.com)) and values 292 were gathered from PubChem (62). The solvents used to prepare stock solutions and mobile phases were: acetonitrile (ACN, HPLC grade) purchased from Fisher Chemical 293 294 (Loughborough, UK); formic acid (99%) purchased from Acros Organics (Geel, Belgium); and 295 ammonium formate (\geq 99.995%) purchased from Sigma-Aldrich (Diegem, Belgium). Ultra-296 pure water (H₂O, conductivity = $0.1 \,\mu$ S/cm, pH 6.00) was produced in the laboratory using a 297 Milli-Q gradient purification system from Millipore (Bedford, MA, USA). 298

- 299 **3.2** Apparatus
- 300

301 All measurements were done on an Agilent Infinity 1290 system from Agilent Technologies 302 (Waldbronn, Germany) that consisted of the following modules: a quaternary pump (G4204A), 303 an autosampler (G4226A), a thermostatted column compartment (G1316C) and a diode array 304 detector with a 1.0 µL flow cell (G4212A). OpenLab software (Agilent Technologies) was 305 used to operate the system and acquire and analyse the data. Further data treatment was done 306 in Microsoft Excel. The maximum pressure of the system was 1200 bar. The injection volume 307 was kept constant at 0.50 μ L and the flow rate was 0.500 mL/min. The detector was set at an 308 acquisition rate of 40 Hz to measure the retention times of the compounds and at 80 Hz to 309 measure the column void and extra-column times. The considered wavelengths were 240 nm, 310 254 nm and 275 nm. The column was placed in an oven with a constant temperature of 30.0 311 °C. Analyses were performed on an Xbridge C_8 column (2.1 x 50 mm; 2.5 µm) from Waters 312 (Wexford, Ireland). Connections between the autosampler, column and detector were made 313 with nanoViper tubing (inner diameter: 75 µm, total length: 1000 mm) from Thermo Scientific 314 (Germering, Germany).

315

316 **3.3 Stock and working solutions**

317

Stock solutions of all compounds were prepared in pure organic (ACN) or aqueous solvent, depending on their solubility, as mentioned for each compound in Table S-1, in a concentration up to 10,000 μ g/mL. Working solutions were made by diluting the stock solutions to 20 μ g/mL in a final solvent mixture of 1.000 mL 5/95 ACN/H₂O.

- 323 **3.4 Dataset**
- 324

Retention factors were collected for 82 representative small molecules (see Table S-1), covering a wide range of physiochemical properties, at ten different isocratic strengths wherein the fraction of ACN (φ_{ACN}) was varied between 0.05 and 0.90 (in 0.10 intervals between 0.10 and 0.90). The aqueous component of the mobile phase was an ammonium formate buffer (brought to pH 3.00 with formic acid). Each compound was injected in triplicate and obtained retention times were averaged. The obtained retention times were converted into retention factors using the following equation:

333
$$k = \frac{t_r - t_0}{t_0 - t_e}$$
 (9)

334

Wherein t_r is the retention time of a particular compound, t_0 is the elution time of an unretained marker (thiourea in this study) and t_e is the extra-column dead time, obtained by replacing the column in the system by a zero dead-volume union and injecting thiourea under the same experimental conditions as for t_r and t_0 .

339

340 Some compounds in the dataset were so strongly retained at low φ_{ACN} (e.g., 0.05 and 0.10) that 341 it became impractical to record their retention factors in a reasonable amount of time. 342 Therefore, the retention factors at low φ_{ACN} for these compounds were estimated by fitting the 343 Neue-Kuss model (Eq. 1) to the available retention factors, and then using the fitted Neue-Kuss 344 model to determine the missing retention factors. Some other compounds were highly polar 345 and did not retain adequately (defined as having a retention factor below 2.5 at 0.05 φ_{ACN} or 346 having less than five retention factor values above 0.001) resulting in unreliable data. These 347 compounds were therefore discarded in this study. The final remaining number of compounds 348 was 57. Furthermore, all datapoints were clipped to have a lowest k-value of 0.001, which was 349 necessary because inherent measurement errors caused low retention factor values to be 350 unreliable, with fluctuations between small positive (<0.001) and small negative (> -0.001) 351 values. All experimentally obtained retention factors are shown in Table S-2 in the supporting 352 information. In addition to isocratic retention data, experimental gradient retention data were 353 additionally collected for the 57 compounds. For this purpose, four different gradient profiles 354 were experimentally run (see Table 1).

- 355 356
- 357 3.5 Compound simulator
- 358

To train the *Q*-network, a compound simulator was created to generate sufficient training data for the *Q*-learning algorithm. By fitting the Neue-Kuss model to all the experimentally obtained retention data for the 57 compounds described above, ranges of parameter values (S₁, S₂ and k_w) were deduced, and these ranges and their mutual relations laid the basis to generate retention parameters for simulated compounds (see Figure S-1 and S-2 in the Supporting Information). The simulator generated a simulated compound in five steps:

- 365
- 366 1. S_I was randomly sampled between $10^{0.9}$ and $10^{1.8}$;
- 367 2. S_2 was sampled from its relationship with S_1 : $S_2 = 2.5010 \cdot \log S_1 2.0822 + r_1$, 368 where r_1 is a random number sampled from a uniform distribution $U \sim (-0.35, 0.35)$:
- 368 where r_1 is a random number sampled from a uniform distribution $U \sim (-0.35, 0.35)$; 369 3. k_w was sampled from its relationship with S_1 : $k_w = 10^{0.0839 \cdot S_1 + 0.5054 + r_2}$, where r_2 is
- a random number sampled from a uniform distribution $U \sim (-1.2, 1.2)$;
- 3714. the resulting parameter values were input to the Neue-Kuss model to output ten372datapoints (retention factors) between $\varphi_{ACN} = 0.05$ and 0.90;
- 3735. random noise sampled from a normal distribution $N \sim (1.0, 0.1)$ was added to each374datapoint via multiplication.
- 375

The last step was included to mimic the noisy nature of the experimental data. A total of 10,000 compounds were simulated according to the five-step procedure above; each of which would occupy a single episode in the complete training of the deep *Q*-learning algorithm. Although not directly evaluated, 10,000 simulated compounds were considered sufficient to capture the compound space for the specific HPLC setup.

385 **3.6 Double deep** *Q***-learning algorithm**

386

388

387 3.6.1 Environment

The dynamics of the reinforcement learning environment aimed to mimic that of a real
chromatographic workflow: sequentially selecting [isocratic] scouting runs for *one* compound
at a time (see Figure 1).

392

393 For consistency, and to allow for evaluation on experimental data later on, the state and action 394 space of the environment was restricted to the fractions of organic modifier (ϕ_{ACN}) that had 395 been experimentally run (namely 0.05 and 0.10 to 0.90 in steps of 0.10; Table S-2). 396 Specifically, a state was defined as an array of 10 elements, each of which was a placeholder 397 for a retention factor value at a certain φ_{ACN} (e.g. the first element was a placeholder for the 398 retention factor value at 0.05 φ_{ACN} , the second element was a placeholder for the retention 399 factor value at 0.10 φ_{ACN} , and so on). The default value of the elements in the array when a 400 given placeholder was not holding any retention factor value was set to -1, as is the case when 401 no scouting runs had been run (see Fig. 1). The value was set to -1 and not 0 in an attempt to 402 make the *Q*-network more easily distinguish between a non-retention factor value and a low 403 retention factor value. Similar to the state space, the action space was defined to select from 404 the different φ_{ACN} that had been run experimentally, with an addition of a STOP action, 405 allowing the agent to stop performing scouting runs (see Fig. 1). At *each* step, the agent was 406 given the choice to select *one* specific scouting run, or to stop.

407

408 To teach the agent to select desirable scouting runs that can be used to accurately model the 409 retention behavior of a given compound, a reward function was defined. The reward function 410 was defined in such a way as to teach, or *reinforce*, the agent to select optimal scouting runs. 411 After the agent had finished performing scouting runs (and at least three different scouting runs 412 had been selected and run), the resulting retention factors (from those three scouting runs) were 413 used to fit the Neue-Kuss model (see appendix A for more information). Subsequently, the 414 reward function computed a reward based on the inverse of the mean relative error (MRE⁻¹) 415 between the target retention factors (y) and the retention factors predicted by the Neue-Kuss 416 model (\hat{y}) :

417

419

418 reward = MRE^{-1} ,

420 where MRE
$$= \frac{1}{10} \sum_{i=1}^{10} \frac{|y_i - \hat{y}_i|}{|y_i|}$$
 (11)

421

422 The targets y are assumed to be the *true* retention factors which we want the predictions \hat{y} to 423 be as close as possible to. In the best-case scenario, true retention factors would be obtained 424 from real experiments. However, as mentioned before, because O-networks need a lot of 425 training data, simulated compounds were used instead; and the relatively few (57) experimental 426 compounds were held-out as a test set, and later used to evaluate the agent after it had been 427 trained. Although the training data were randomly sampled using the simulator, which used 428 ranges of Neue-Kuss parameter values $(S_1, S_2 \text{ and } k_w)$ that were based on those of the 429 compounds in the held-out test set, these parameters were merely roughly deduced from the 430 overall trends of the parameter values of the test compounds and should therefore not 431 undermine the purpose of the held-out test set, which is to evaluate the agent on real 'unseen' 432 data. Importantly, the MRE was always calculated based on all (10) retention factors (0.05 \rightarrow 433 0.90 φ_{ACN}), as the purpose was to obtain models that predicted well for all (10) φ_{ACN} .

(10)

Furthermore, if the agent stopped before three scouting runs had been selected, the MRE
evaluation (Eq. 10) was not carried out and the agent was rewarded zero (0). Finally, the agent
was penalized (negatively rewarded) in two ways:

- 437
- 438 439

1. by selecting scouting runs that had already been selected before, penalized -5; 2. by selecting scouting runs that resulted in long analysis times (> 1 hour), penalized based on a sigmoidal (s-shaped) function f_s :

440 441 442

443

$$f_s(k) = \frac{1}{1 - e^{-(k \cdot 0.00425 - 4.0)}} \times 20 \tag{12}$$

444 where k is the retention factor. The function f_s is adapted in such a way as to have a maximum 445 penalty close to the maximum reward that can be obtained (Eq. 10), and a minimum penalty at 446 zero (0). These penalties were implemented to enable the agent to quickly learn not to select 447 the same scouting run multiple times, and to avoid selecting scouting runs that would result in 448 high analysis times, respectively. The analysis time penalty (Eq. 12) was important because the 449 Q-network had to learn to tailor the choice of scouting runs depending on the retention behavior 450 of the compound. For example, if the agent had to select scouting runs for a highly retained 451 compound, it was expected to avoid running scouting runs at low φ_{ACN} . 452

453 3.6.2 Agent

In this study, as an extension to the deep *Q*-learning algorithm introduced in section 2, a *double deep Q-learning algorithm* (63) with *experience replay* (64,65) was implemented.
Additionally, an *epsilon-greedy policy* (60) was incorporated for selecting actions (see appendix A, B and C for details on these implementations respectively).

459

460 Instead of having a single *Q*-network for both the target and the prediction together, the double 461 deep *Q*-learning algorithm utilizes two separate neural networks with two separate *Q*-functions 462 to approximate Q^* : a *target* network and a *prediction* network, for the target and the prediction 463 respectively. The minimization procedure is based on the procedure of a single *Q*-network as 464 described in (Eq 8). The target network has fixed parameters θ^- which are updated every *N* 465 episodes by having the prediction network copying over its parameters θ , while the prediction 466 network is updated every episode via the minimization procedure (Eq. 8).

467

468 It is of great importance to have a separate network to produce the target, because it counters 469 the issue of overestimating and biasing the target. The overestimation (and bias) of the target 470 naturally occurs for standard *Q*-learning and DQN due to the max operator (cfr. max_a in eq. 5 471 and 6) both selecting and evaluating the actions. Hasselt et al. have shown how Double deep 472 *Q*-learning produces more accurate estimations of the expected return as well as better policies 473 (63,66).

474

475 A crucial quality of the minimization procedure (Eq. 8) is that it can be formulated to accept 476 diverse batches of non-correlated examples. Specifically, the minimization step can be done 477 batch-wise, where each batch contains transitions from different trajectories. In other words, each batch may contain tuples (s,a,s') from entirely different scouting runs. To be able to 478 479 sample such batches, a memory unit called *replay memory* was implemented to store transitions 480 that were experienced by the agent while it was performing scouting runs. These stored 481 transitions were scheduled to be sampled (in batches) after each episode, and to be directly 482 used to train the agent (minimize the objective function). Although we did not test any

483 algorithm without experience replay (replay memory), it is believed that this technique greatly 484 stabilizes and smoothens the training process (64,65). The discount factor γ was set to 0.95 as 485 an attempt to give slightly higher weight to early rewards.

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489

488 **3.7** Evaluation

490 After the double deep *Q*-learning agent was trained, it was evaluated on real, experimental data. 491 Similar to how the agent selected scouting runs for simulated data during training, the agent 492 selected scouting runs for isocratic experimental compounds. The resulting Neue-Kuss model 493 (fitted on the retention factors resulting from the scouting runs selected by the agent) were then 494 used to predict the retention factors for all 10 isocratic scouting runs. Additionally, the same 495 Neue-Kuss parameters were also used in the Neue-Kuss equation for gradient elution, which 496 was used to predict retention data under four different gradient conditions (Table 1). The model 497 for gradient elution is defined as follows:

498

499
$$t_{r,gradient} = \frac{1}{S\beta} \frac{(1+S_2\varphi_0)^2 \ln(1+S_1\beta k_w \exp(\frac{-S_1\varphi_0}{1+S_2\varphi_0})(t_0-\frac{t_D}{k_0}))}{1-S_2(\frac{1+S_2\varphi_0}{S_1}) \ln(1+S_1\beta k_w \exp(\frac{-S_1\varphi_0}{1+S_2\varphi_0})(t_0-\frac{t_D}{k_0}))} + t_0 + t_D$$
(13)

500

501 where S₁, S₂ and k_w are the same parameters as for the isocratic model (cfr Eq. 1), t_{r,gradient} 502 is the gradient retention time, φ_0 the fraction of strong eluting solvent in the mobile phase at 503 the start of the gradient, β the slope of the gradient (= $\Delta \phi/t_G$), k_0 the retention factor at ϕ_0 , t_0 504 the elution time of an unretained compound and t_D the dwell time of the system (the time the mobile phase needs to flow from the pump to the head of the column). The predicted retention 505 506 factors (for both isocratic and gradient runs) were then compared to all experimental data 507 available for that compound by calculating the mean relative percentage error (MRPE) with 508 modification, as follows:

509

$$MRPE = \frac{1}{10} \sum_{i=1}^{10} \frac{|y_i - \hat{y}_i|}{|1 + y_i|} \times 100$$

511

The (modified) MRPE was used as it was less sensitive to small k-values (<0.1), creating a more stable and robust evaluation metric. To further illustrate how well the resulting model (based on the selected scouting runs) performed, a comparison was made to a model fit on all available isocratic datapoints ($0.05 \rightarrow 0.90 \ \varphi_{ACN}$), a model fit on a random selection of three isocratic scouting runs, as well as a model fit on a chromatographer's selection of three scouting runs, namely $\varphi_{ACN}= 0.1, 0.5$ and 0.8.

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(14)

521 4 RESULTS AND DISCUSSION

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524

523 4.1 Learning curves: training progression

525 To illustrate how the agent learned to perform optimal scouting run selection, the *rewards* and 526 number of selected scouting runs were collected during the training procedure. Figure 2 527 illustrates both how the reward and the number of selected actions changed over the first 3500 528 episodes. In addition, for comparison, the theoretical optimum (highest possible reward) was 529 also calculated and plotted. Notice that the theoretical optimum does not reflect the best 530 possible performance by an agent, but rather illustrates what in theory could be obtained. 531 Specifically, the theoretical optimum was obtained by selecting, for each and every compound 532 separately, the combination of (three) scouting runs (out of the 120 combinations that exist for 533 3 actions and 10 possible states) that resulted in the highest reward. Due to random action 534 selection, leading to poor retention models and high penalties, the reward given to the agent 535 started off at high negative values (around -5 to -15). Specifically, the low reward as well as 536 the high number of selected scouting runs in the first 100 episodes confirmed that the selected 537 actions were highly random (due to the high ϵ for the epsilon-greedy policy). Between episode 538 100 and 200 however, the agent started to select scouting runs non-randomly, which resulted 539 in the agent selecting much fewer scouting runs (around 2 on average). Notice that, although 540 not yielding a retention model due to too few scouting runs (\leq 3), the agent had at this point 541 learned to select better actions than before, because no or fewer penalties were given. After 542 200 episodes, the agent went from an insufficient number of scouting runs (which resulted in 543 a reward around 0) to selecting enough scouting runs for a retention model, resulting in a 544 positive reward outweighing potential penalties. In other words, after 200 episodes, the agent 545 started to learn how to optimize the selection of scouting runs for retention modelling -546 selecting around four scouting runs on average, which is on average one scouting run more 547 than the minimum required to fit a three-parameter model. This number was overestimated due 548 to the epsilon-greedy policy ($\epsilon > 0$), which is likely to cause the agent to select scouting runs 549 sub-optimally (i.e. not selecting actions with the greatest <u>*Q*-value</u>). Notice, although not 550 visualized, if only the greedy policy was followed (epsilon = 0), the agent would select closer 551 to three scouting runs on average.

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554

4.2 Scouting run selection

555 After the agent, or the *Q*-network, was optimized or trained to select scouting runs for simulated 556 data, it was evaluated on experimental data. Figure 3 illustrates how the agent selected scouting 557 runs for four representative compounds it had never seen before (see Table S-3 for complete 558 results). For the four compounds considered in Figure 3, φ_{ACN} of 0.20 and 0.90 were selected 559 by the trained agent; and for all four compounds, three scouting runs were run in total (which 560 is the minimum requirement). What differed between compounds of different retention 561 behavior, specifically between less retained compounds (e.g. nitrobenzene and o-562 methylacetophenone) and more retained compounds (e.g. iodobenzene and anthracene), was the selection of the third scouting run (φ_{ACN} of 0.05 for the less retained compounds and 0.50 563 564 for the more retained compounds), which indicates that the selection of scouting runs was 565 tailored based on the retention behavior of the compounds. Because nitrobenzene and o-566 methylacetophenone are less retained compounds, they were both selected to be run at as low 567 as $\varphi_{ACN}=0.05$. Intuitively, a scouting run at $\varphi_{ACN}=0.05$ holds important information for a 568 retention model but requires significantly longer analysis time for highly apolar compounds, 569 like iodobenzene and anthracene. For these highly retained compounds, $\varphi_{ACN}=0.05$ was avoided and instead $\varphi_{ACN}=0.50$ was selected by the agent. Furthermore, in all cases, the scouting run 570

selection spanned a wide range of selectable φ_{ACN} , in a relatively evenly spaced-out manner, which intuitively should result in more accurate retention models. Finally, these results illustrate that the penalty given to the agent for selecting scouting runs resulting in high (>> 1 hour) analysis times (i.e. high retention factors), made the agent avoid the lower percentages for compounds like iodobenzene and anthracene, but not for compounds like nitrobenzene and o-methylacetophenone, which in regards to this study was highly desirable.

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- 578

579 4.3 Q-values: taking action

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581 While Figure 3 illustrates which actions the agent took depending on the retention behavior of 582 the compounds, Figure 4 illustrates how the agent, or the Q-network and its Q-values, decided 583 which actions to take, given a certain state. It also illustrates how the O-values varied between 584 steps and between compounds with a different retention behavior (specifically, in Figure 4, 585 acetophenone and biphenyl). Interestingly, the penalty given for high analysis times, forced the 586 *Q*-values for $\varphi_{ACN} = 0.05$ and $\varphi_{ACN} = 0.10$ to differ significantly between the two compounds. Specifically, biphenyl, which is more retained than acetophenone, was predicted by the Q-587 588 network to result in a high penalty if run at $\varphi_{ACN}=0.05$ or $\varphi_{ACN}=0.10$ (the analysis time would be too long), and therefore predicted higher Q-values at higher φ_{ACN} . Furthermore, the penalty 589 given when the same φ_{ACN} was selected more than once, resulted in a significant lowering of 590 591 the *Q*-value in the succeeding steps for that φ_{ACN} .

592

Acetophenone had higher *Q*-values on average, suggesting the reward/penalty on average was greater for less retained compounds (like acetophenone) than higher retained compounds (like biphenyl). This suggests that (1) $\varphi_{ACN}=0.05$ was an important datapoint to accurately model retention behaviors (resulting in a low MRPE) – a datapoint which did not get selected for highly retained compounds due to high penalties; (2) the continual higher penalties for highly retained compounds forced down the *Q*-values.

599

Although not directly presented in this study, the *number* of selected scouting runs, as well as the specific selection of scouting runs may differ between agents trained on slightly different data. Specifically, Figure 4 illustrates how several actions have similar Q-values given a certain state, which could easily nudge the agent in a different direction. This does not necessarily mean that the agent (or the *double deep Q-learning algorithm*) is not robust, but rather that there are several solutions to the problem – *i.e.* several directions (combinations of scouting runs) that result in good retention models.

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4.4 Retention model performance: isocratic and gradient prediction errors

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To get a better understanding of how well the retention models resulting from the scouting runs selected by the agent perform, the prediction accuracy of these retention models was compared to the prediction accuracy of retention models obtained by fitting to (1) all 10 available isocratic data points (φ_{ACN} of 0.05 to 0.90); (2) a random selection of three isocratic scouting runs and (3) a chromatographer's selection of three isocratic scouting runs, namely φ_{ACN} = 0.1, 0.5 and 0.8. These four differently obtained retention models were also used to predict gradient retention factors for four different gradients (Table 2).

617

Table 2 illustrates the MRPE between the experimentally obtained data and the predictions, forboth isocratic and gradient data (see Table S-4 for complete results). The retention model

620 obtained by the agent (based on (mostly) 3 scouting runs), performed comparably and better to 621 the retention models based on all experimental datapoints and the retention models based on 622 the chromatographer's selection of three scouting runs, for the isocratic and gradient data 623 respectively; and performed significantly better than the retention models based on the random 624 scouting run selection. Importantly, the models resulting from the agent's selection of scouting 625 runs were, compared to retention models based on all datapoints, obtained via seven fewer 626 datapoints; *i.e.* seven fewer scouting runs, saving significant time and costs. These results 627 indicate that the agent successfully learned to optimize the scouting run selection.

628

629 The reason why the retention model obtained by the agent had a significantly lower gradient 630 prediction error compared to the retention model fitted on all datapoints and the 631 chromatographer's selection, lies in the high prediction error for the lowly retained compounds. 632 It is speculated that (1) having many datapoints at medium-high φ_{ACN} for the lowly retained compounds, where most φ_{ACN} have retention factors close to 0, is forcing the estimated 633 634 parameters (k_w, S₁ and S₂) to model the behavior of the given lowly retained compound favorably in the regions of low φ_{ACN} , but highly unfavorably in the regions of high φ_{ACN} , which 635 636 is more relevant for gradient retention modelling; and (2) not having φ_{ACN} of 0.05 (in the case 637 of the chromatographer's selection) misses out on valuable information for gradient 638 predictions.

640 5 Conclusions

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639

642 In this study, a reinforcement learning algorithm, specifically the double deep O-learning 643 algorithm, was shown to be able to learn to optimally select informative scouting runs in a fully 644 self-taught way. Although only isocratic scouting runs were considered for a specific RPLC 645 setup, these experiments illustrate how the agent learned to tailor the selection of scouting runs 646 for different compounds depending on the retention behavior (mainly defined by its polarity, 647 or k_w). The experiments also illustrate how the agent limited the number of selected scouting 648 runs yet still yielding a retention model with low prediction error (MRPE of 3.77% and 1.93%) 649 for isocratic and gradient data, respectively). The strategy of the agent was to select relatively 650 evenly spaced-out scouting runs (in terms of φ_{ACN}), including at least one scouting run at low 651 φ_{ACN} (as long as the analysis time was not too long), and at least one scouting run at high φ_{ACN} . 652 Intuitively, selecting scouting runs as such will cover a greater space, better capturing the 653 complete behavior of the compound. The retention models based on the agent's selection of 654 scouting runs (MRPE of 3.77% for the isocratic data and 1.93% for the gradient data) compared 655 well to both the retention models fitted on all datapoints (MRPE of 3.26% for isocratic data 656 and 4.97% for gradient data) and the chromatographer's selection (3.86% for isocratic data and 657 6.66% for gradient data); and performed significantly better than retention models based on the 658 random selection of three scouting runs (MRPE of 46.30% for isocratic data 7.60% for gradient 659 data).

660

661 Although the double deep *O*-learning algorithm presented in this study shows potential, it has 662 only been tested on its ability to learn from isocratic data for a specific RPLC setup. This double 663 deep *Q*-learning algorithm is also limited to a discrete, and preferably small, action space, 664 which could potentially be a problem for tasks like selecting scouting runs for gradient elution 665 - where the action space is either larger or continuous. The prospect is to develop a reinforcement learning algorithm that can deal with a more complex/larger state and action 666 667 space (e.g. Branching Dueling Q-learning (67)) and/or continuous action space (e.g. Twin-668 Delayed Deep Deterministic Policy Gradient (TD3) or Soft Actor-Critic (SAC) algorithm (68,69)), to be able to perform scouting runs for a mixture of compounds (*i.e.* selecting scouting 669

670 runs for multiple compounds at the same time) in a more complex setting such as gradient 671 conditions. If such an algorithm can be developed successfully, it would be worthwhile 672 integrating it into normal practice.

673 674

675 Availability

All code used in this study (except the plots), including the complete implementation of the 676 677 agent and the environment, can be found at https://github.com/akensert/ddqn-isocraticscouting-runs. Due to stochasticity in training an artificial neural network, results may differ 678 679 somewhat from run to run. InChI for each experimental compound evaluated in this study can 680 be found in supplementary table 5 (Table S-5).

681

682 **Conflicts of interest**

There are no conflicts of interest. 683

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693 **Credit Author Statement**

- 694 Alexander Kensert: Conceptualization; Data curation; Formal analysis; Investigation;
- 695 Methodology; Validation; Visualization; Writing - original draft.
- 696 Gilles Collaerts: Data curation; Formal analysis; Investigation; Validation; Writing - review 697 & editing.
- 698 Kyriakos Efthymiadis: Formal analysis; Investigation; Methodology; Writing – review & 699 editing.
- 700 Gert Desmet: Conceptualization; Funding acquisition; Investigation; Methodology;
- 701 Supervision; Writing - review & editing.
- 702 Deirdre Cabooter: Conceptualization; Funding acquisition; Formal analysis; Methodology;
- Investigation; Supervision; Writing review & editing. 703

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915 Figure Captions

916

Figure 1: A simple schematic illustration of the interaction between agent and environment. A) the environment supplies the agent with a state *s*, which is an array of retention factors (k) where each element corresponds to a φ_{ACN} ; based on this state *s* the agent produces an array of *Q*-values for each action. B) the agent takes the greedy action, namely the action with the highest *Q*-value, resulting in a new state *s'*. For readability, the figure excludes the reward and only covers one single transition (s,a,s').

923

Figure 2: Illustration of how the reward (top; blue) and the number of scouting runs selected (bottom; orange) by the agent increased over the number of episodes. To reduce the noise of these learning curves, a moving average filter was applied to the data (window size of 35). The lighter shaded areas represent a 95% confidence interval of the average lines. The black line in the top plot illustrates the theoretical optimum (the highest possible reward) that can be obtained by selecting the best combination of three scouting runs out of 120 different combinations.

931 Figure 3: Illustration of how the scouting runs were selected sequentially for four 932 representative compounds with a significantly different retention behavior. The estimated 933 Neue-Kuss parameter k_w can be used to assess the retention behavior of the compounds, with 934 a high k_w indicating strong retention, and a low k_w indicating weaker retention. The blue dots 935 indicate the experimental data, the orange triangles pointing to the x-axis indicate the scouting 936 runs selected (sequentially, $1 \rightarrow 3$), and the orange lines indicate the predicted retention model 937 resulting from the scouting runs. MRPE stands for mean relative percentage error. The y-axis 938 has been set to log scale to better visualize the prediction of lower φ_{ACN} values.

939

940Figure 4: Illustration of how the scouting runs were sequentially (from top to bottom) selected941based on the highest Q-value at each step, for two representative compounds. Similar to Figure9423, the estimated Neue-Kuss parameter k_w indicates the retainability of the compounds, with a943high k_w indicating strong retention, and a low k_w indicating weaker retention. Orange bars944

- 945
- 946
- 947

Appendix:

Deep Q-learning for the selection of optimal

950 isocratic scouting runs in liquid chromatography

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948

952 A: Retention Model

953 The retention model used in this study was the Neue-Kuss model (Eq. 1). The model is obtained 954 by *fitting* the model's parameters $(S_1, S_2 \text{ and } k_w)$ to the available datapoints for a given 955 The fitting done via Nelder-Mead optimization (doi: compound. is 956 https://doi.org/10.1093/comjnl/7.4.308), which is a type of simplex method.

957

958 **B: Double Deep Q-network**

959 In this study, the two Q-networks had a 10-unit input layer (corresponding to the state of the 960 environment), two 1024-unit hidden layers with rectified linear unit activation and a dropout 961 rate of 0.2, and a final 11-unit linear output layer (corresponding to the actions of the agent). 962 While, the prediction network was updated each episode via the minimization procedure, the 963 target network was updated (by copying over the prediction network's parameters) every 64 964 steps. The batch-size, which is the number of transitions used per minimization step (Eq. 8), 965 was set to 128. To minimize the objective function (Eq. 8), a stochastic gradient descent 966 optimizer was used, with a momentum and an initial learning rate of 0.9 and 0.001 respectively. 967 The learning rate was scheduled to decay for 4096 iterations (episodes) until it reached a 968 minimum learning rate of 0.0001.

969

970 Before selecting the hyperparameter values mentioned above, some preliminary testing was 971 done (i.e. the Q-network was tested with different hyperparameter values). The number of 972 hidden units was varied between 512 and 2048 (for which 1024 was equal or better to the other 973 values and was therefore selected), the number of layers were varied between 1 and 3 (for 974 which 2 was selected), and the dropout rate was varied between 0.1 and 0.5 (for which 0.2 was 975 selected). Furthermore, the initial learning rate was varied between 0.01 and 0.001 (for which 976 0.001 was selected), and both rectified linear unit (ReLU) and sigmoid activation was tested 977 (for which ReLU was selected). Finally, discount factor γ was varied between 0.0 and 1.0 (for 978 which 0.95 was selected).

979

980 C: Replay Memory/Experience Replay

981 Because the batch-size was set to 128, the number of samples sampled from the replay memory 982 unit was also 128. The capacity (maximum number of transitions stored) for the replay memory 983 unit was restricted to 2048. This restriction was added to avoid having the Q-network learn 984 from old experiences (transitions). The sampling of transitions from the replay memory unit 985 did not start until it had at least 512 transitions stored; because the Q-network (or specifically 986 the prediction network) was updated every episode, having at least 4 times the batch-size of 987 transitions helped to avoid having the Q-network train on similar batches multiple times in a 988 row; and although no evidence is presented here, it was thought that this could potentially bias 989 the O-network at the beginning of training and hence slow down learning. As for the 990 hyperparameters of the Q-network the batch size was varied between 32 and 512 (for which 991 128 was selected).

993 **D: Epsilon-greedy policy**

994 The epsilon-greedy policy is important for balancing the exploration-exploitation tradeoff. The epsilon-greedy policy in this study was divided into $\varepsilon_{initial}$, ε_{decay} , and $\varepsilon_{minimum}$, which 995 were set to 1.0, 0.99, and 0.1 respectively. This made the agent take mostly random actions 996 997 (exploring) in the beginning, and increasingly by time started to take actions based on the 998 output of the Q-function (the Q-values), namely $\arg \max_a Q(s, a)$ (exploiting), The minimum 999 epsilon was kept at 0.1 to force the agent to explore throughout the entire run. As for the 1000 hyperparameters of the *Q*-network, the initial epsilon, the decay rate, as well as the minimum 1001 decay was varied between 0.1-1.0, 0.99-0.999 and 0.0-1.0 respectively. For which 1.0, 0.99 1002 and 0.1 was selected respectively. 1003