The concentrations of measure phenomena were discovered as the mathematical background to statistical mechanics at the end of the nineteenth/beginning of the twentieth century and have been explored in mathematics ever since. At the beginning of the twenty-first century, it became clear that the proper utilization of these phenomena in machine learning might transform the curse of dimensionality into the blessing of dimensionality. This paper summarizes recently discovered phenomena of measure concentration which drastically simplify some machine learning problems in high dimension, and allow us to correct legacy artificial intelligence systems. The classical concentration of measure theorems state that i.i.d. random points are concentrated in a thin layer near a surface (a sphere or equators of a sphere, an average or median-level set of energy or another Lipschitz function, etc.). The new stochastic separation theorems describe the thin structure of these thin layers: the random points are not only concentrated in a thin layer but are all linearly separable from the rest of the set, even for exponentially large random sets. The linear functionals for separation of points can be selected in the form of the linear Fisher’s discriminant. All artificial intelligence systems make errors. Non-destructive correction requires separation of the situations (samples) with errors from the samples corresponding to correct behaviour by a simple and robust classifier. The stochastic separation theorems provide us with such classifiers and determine a non-iterative (one-shot) procedure for their construction.
This article is part of the theme issue ‘Hilbert’s sixth problem’.

1. Introduction: five ‘foundations’, from geometry to probability, quantum mechanics, statistical physics and machine learning

It’s not given us to foretell
How our words will echo through the ages,…

F. I. Tyutchev, English Translation by F. Jude

The sixth Hilbert problem was inspired by the ‘investigations on the foundations of geometry’ [1], i.e. by Hilbert’s work *The foundations of geometry* [2], which firmly implanted the axiomatic method not only in the field of geometry but also in other branches of mathematics. The sixth problem proclaimed expansion of the axiomatic method beyond existing mathematical disciplines, into physics and further on.

The sixth problem sounds very unusual and not purely mathematical. This may be a reason why some great works which have been inspired by this problem have no reference to it. The most famous example is the von Neumann book [3] *Mathematical foundations of quantum mechanics*. John von Neumann was an assistant to Hilbert, and they worked together on the mathematical foundation of quantum mechanics. This work was obviously in the framework of the sixth problem, but this framework was not mentioned in the book.

In 1933, Kolmogorov answered the Hilbert challenge of axiomatization of the theory of probability [4]. He did not cite the sixth problem but explicitly referred to Hilbert’s *Foundations of geometry* as the prototype for ‘the purely mathematical development’ of the theory. But Hilbert in his sixth problem asked for more, for ‘a rigorous and satisfactory development of the method of the mean values in mathematical physics’. He had in mind statistical physics and ‘in particular the kinetic theory of gases’. The sixth chapter of Kolmogorov’s book contains a survey of some results obtained by Kolmogorov and Khinchin about independence and the law of large numbers, and the appendix includes a description of the 0-1 laws in probability. These are the first steps to a rigorous basis of ‘the method of mean values’. Ten years later, in 1943, Khinchin published a book entitled *Mathematical foundations of statistical mechanics* [5]. This brought an answer to the sixth problem one step closer, but again without explicit reference to Hilbert’s talk. The analogy between the titles of von Neumann’s and Khinchin’s books is obvious.

The main idea of statistical mechanics, in essence, can be called the *blessing of dimensionality*: if a system can be presented as a union of many weakly interacting subsystems, then, in the thermodynamic limit (when the number of such subsystems tends to infinity), the whole system can be described by relatively simple deterministic relations in the low-dimensional space of macroscopic variables. *More means less*—in very high-dimensional spaces many differences between sets and functions become negligible (vanish) and the laws become simpler. This point of view on statistical mechanics was developed mainly by Gibbs [6] (ensemble equivalence), but Khinchin made the following remark about this work: ‘although the arguments are clear from the logical standpoint, they do not pretend to any analytical rigor’ [5, p. 4], exactly in the spirit of Hilbert’s request for ‘a rigorous and satisfactory development’. The devil is in the detail: how should we define the thermodynamic limit and in which sense are the ensembles equivalent? For some rigorously formulated conditions, the physical statements become exact theorems.

Khinchin considered two types of background theorems: ergodic theorems and limit theorems for high-dimensional distributions. He claimed that the foundations of statistical mechanics should be a complete abstraction from the nature of the forces. Limit theorems use very general
properties of distributions in high dimension, indeed, but the expectations that ergodicity is a
typical and universal property of smooth high-dimensional multi-particle Hamiltonian systems
were not met [7]. To stress that the ergodicity problem is non-trivial, we have to refer to
the Oxtoby–Ulam theorem about metric transitivity of a generic continuous transformation,
which preserves volume [8] (we see that typical properties of continuous transformations differ
significantly from typical properties of smooth transformations).

Various programmes proposed for the mathematical foundation of statistical mechanics were
discussed, for example, by Dobrushin [9] and Batterman [10]. Despite the impressive proof
of ergodicity of some systems (hyperbolic flows or some billiard systems, for example), the
now became dominant; the ergodic hypothesis is neither a necessary nor a sufficient condition
for the foundation of statistical mechanics (Dobrushin [9] attributed this opinion to Lebowitz,
while Jaynes referred to Gibbs [6], who, perhaps, ‘did not consider ergodicity as relevant to the
foundation of the subject’ [11, p. 87]).

Through the efforts of many mathematicians, the limit theorems from probability theory and
results about ensemble equivalence from the foundation of statistical physics were developed
far enough to become the general theory of measure concentration phenomena. Three articles
were especially important for our work [12–14]. Ledoux [15] gives an introduction into the
mathematical theory of measure concentration. A simple geometric introduction into these
phenomena was given by Ball [16].

Perhaps the simplest manifestation of measure concentration is the concentration of the
volume of the high-dimensional ball near the sphere. Let \( V_n(r) \) be a volume of the \( n \)-dimensional
ball of radius \( r \). It is useful to stress that the ‘ball’ here is not necessarily Euclidean and means
the ball of any norm. Lévy [17] recognized this phenomenon as a very important property of
geometry of high-dimensional spaces. He also proved that equidistributions in the balls are
asymptotically equivalent in high dimensions to the Gaussian distributions with the same mean
value of squared radius. Gibbs de facto used these properties for sublevel sets of energy to
demonstrate equivalence of ensembles (microcanonical distribution on the surface of constant
energy and canonical distribution in the phase space with the same mean energy).

Maxwell used the concentration of measure phenomenon in the following settings. Consider
a rotationally symmetric probability distribution on the \( n \)-dimensional unit sphere. Then its
orthogonal projection on a line will be a Gaussian distribution with small variance \( 1/n \) (for large
\( n \) with high accuracy). This is exactly the Maxwellian distribution for one degree of freedom in
a gas (and the distribution on the unit sphere is the microcanonical distribution of the kinetic
energy of gas, when the potential energy is negligibly small). Geometrically, it means that if we
look at the one-dimensional projections of the unit sphere, then the ‘observable diameter’ will be
small, of the order of \( 1/\sqrt{n} \).

Lévy noticed that instead of orthogonal projections on a straight line we can use any \( \eta \)-Lipschitz function \( f \) (with \( \|f(x) - f(y)\| \leq \eta \|x - y\| \)). Let points \( x \) be distributed on a unit
\( n \)-dimensional sphere with rotationally symmetric probability distribution. Then the values of
\( f \) will be distributed ‘not more widely’ than a normal distribution around the mean value \( \mathbb{E}_f \); for
all \( \varepsilon > 0 \)

\[
P(\|f - \mathbb{E}_f\| \geq \varepsilon) \leq 2 \exp \left( -\frac{n \varepsilon^2}{2c \eta^2} \right),
\]

where \( c \) is a constant, \( c \leq 9 \pi^3 \). Interestingly, if we use in this inequality the median value of \( f \), \( M_f \),
instead of the mean, then the estimate of the constant \( c \) can be decreased: \( c \leq 1 \). From the statistical
mechanics point of view, this Lévy lemma describes the upper limit of fluctuations in gas for
an arbitrary observable quantity \( f \). The only condition is the sufficient regularity of \( f \) (Lipschitz
property).

Hilbert’s sixth problem influenced this stream of research either directly (Kolmogorov and,
perhaps, Khinchin among others) or indirectly, through the directly affected works. And it
continues to spread this influence to other areas, including high-dimensional data analysis, data mining and machine learning.

At the turn of the millennium, Donoho [18] gave a lecture about the main problems of high-dimensional data analysis with the impressive subtitle: ‘The curses and blessings of dimensionality’. He used the term curse of dimensionality ‘to refer to the apparent intractability of systematically searching through a high-dimensional space, the apparent intractability of accurately approximating a general high-dimensional function, the apparent intractability of integrating a high-dimensional function’ [18, p. 1]. To describe the blessing of dimensionality, he referred to the concentration of measure phenomenon, ‘which suggest that statements about very high-dimensional settings may be made where moderate dimensions would be too complicated’ [18, p. 1]. Anderson et al. [19] characterized some manifestations of this phenomenon as ‘The more, the merrier’.

In 1997, Kainen [20] described the phenomenon of blessing of dimensionality, illustrated it with numerous examples and suggested connections with geometric phenomena in high-dimensional spaces.

The claim of Donoho’s talk was similar to Hilbert’s talk and he cited this talk explicitly: ‘My personal research experiences, cited above, convince me of Hilbert’s position, as a long run proposition, operating on the scale of centuries rather than decades’ [18, p. 28]. The role of Hilbert’s sixth problem in the analysis of the curse and blessing of dimensionality was not mentioned again.

The blessing of dimensionality and the curse of dimensionality are two sides of the same coin. For example, the typical property of a random finite set in a high-dimensional space is: the squared distance of these points to a selected point are, with high probability, close to the average (or median) squared distance. This property drastically simplifies the expected geometry of data (blessing) [21,22], but, at the same time, makes the similarity search in high dimensions difficult and even useless (curse) [23].

Extension of the sixth Hilbert problem to data mining and machine learning is a challenging task. There exists no unified general definition of machine learning. Most classical texts consider machine learning through formalization and analysis of a set of standardized tasks [24–26]. Traditionally, these tasks are:

— classification—learning to predict a categorical attribute using values of given attributes on the basis of given examples (supervised learning);
— regression—learning to predict numerical attributes using values of given attributes on the basis of given examples (supervised learning);
— clustering—joining of similar objects in several clusters (unsupervised learning) [27];
— various data approximation and reduction problems: linear and nonlinear principal components [28], principal graphs [29], independent components [30], etc. (clustering can also be considered as a data approximation problem [31]);
— probability distribution estimation.

For example, Cucker & Smale [24] considered the least squares regression problem. This is the problem of the best approximation of an unknown function \( f : X \rightarrow Y \) from a random sample of pairs \((x, y) \in X \times Y\). Selection of ‘the best’ regression function means minimization of the mean square error deviation of the observed \( y \) from the value \( f(x) \). They use the concentration inequalities to evaluate the probability that the approximation has a given accuracy.

It is important to mention that the Cucker–Smale approach was inspired in particular by J. von Neumann: ‘We try to write in the spirit of H. Weyl and J. von Neumann’s contributions to the foundations of quantum mechanics’ [24, p. 4]. The J. von Neumann book [3] was one step in the realization of Hilbert’s sixth problem programme, as we know. Therefore, the Cucker–Smale ‘mathematical foundation of learning’ is a grandchild of the sixth problem. This is the fourth ‘foundation’ (after Kolmogorov, von Neumann and Khinchin). Indeed, it was an attempt to give a rigorous development of what they ‘have found to be the central ideas of learning
theory’ [24, p. 4]. This problem statement follows Hilbert’s request for ‘rigorous and satisfactory
development of the method of mean values’ [1, p. 454], but this time the development was done
for machine learning instead of mathematical physics.

Cucker and Smale followed Gauss and proved that the least squares solution enjoys
remarkable statistical properties, i.e. it provides the minimum variance estimate [24]. Nevertheless,
non-quadratic functionals are employed as solutions to many problems: to enhance robustness, to
avoid oversensitivity to outliers, to find sparse regression with exclusion of non-necessary input
variables, etc. [25,26]. Even non-convex quasinorms and their tropical approximations are used
efficiently to provide sparse and robust learning results [32]. Vapnik [26] defined a formalized
fragment of machine learning using minimization of a risk functional that is the mathematical
expectation of a general loss function.

Gromov [33] proposed a radically different concept of ergosystems which function by building
their ‘internal structure’ out of the ‘raw structures’ in the incoming flows of signals. The essential
mechanism of ergosystem learning is goal free and independent of any reinforcement. In a
broad sense, loosely speaking, in this concept ‘structure’ = ‘interesting structure’ and learning
of structure is goal free and should be considered as a structurally interesting process.

There are many other approaches and algorithms in machine learning that use some
specific ideas from statistical mechanics: annealing, spin glasses, etc. (see, for example, [34])
and randomization. It was demonstrated recently that the assignment of random parameters
should be data dependent to provide the efficient and universal approximation property of
the randomized learner model [35]. Various methods for evaluation of the output weights
of the hidden nodes after random generation of new nodes were also tested [35]. Swarm
optimization methods for learning with random regeneration of the swarm (virtual particles)
after several epochs of learning were developed in 1990 [36]. Sequential Monte Carlo methods
for learning neural networks were elaborated and tested [37]. A comprehensive overview of the
classical algorithms and modern achievements in stochastic approaches to neural networks was
performed by Scardapane & Wang [38].

In our paper, we do not discuss these ideas; instead, we focus on a deep and general similarity
between high-dimensional problems in learning and statistical physics. We summarize some
phenomena of measure concentration which drastically affect machine learning problems in high
dimension.

2. Waist concentration and random bases in machine learning

After the classical works of Fisher [39] and Rosenblatt [40], linear classifiers have been considered
as the inception of data analytics and machine learning (e.g. [26,41,42], and references therein).
The mathematical machinery powering these developments is based on the concept of linear
separability.

Definition 2.1. Let \( \mathcal{X} \) and \( \mathcal{Y} \) be subsets of \( \mathbb{R}^n \). Recall that a linear functional \( l \) on \( \mathbb{R}^n \) separates
\( \mathcal{X} \) and \( \mathcal{Y} \) if there exists a \( t \in \mathbb{R} \) such that

\[
l(x) > t > l(y) \quad \forall x \in \mathcal{X}, \quad y \in \mathcal{Y}.
\]

A set \( S \subset \mathbb{R}^n \) is linearly separable if, for each \( x \in S \), there exists a linear functional \( l \) such that
\( l(x) > l(y) \) for all \( y \in S, y \neq x \).

If \( \mathcal{X} \subset \mathbb{R}^n \) is a set of measurements or data samples that are labelled as ‘class 1’, and \( \mathcal{Y} \) is a set
of data labelled as ‘class 2’, then a functional \( l \) separating \( \mathcal{X} \) and \( \mathcal{Y} \) is the corresponding linear
classifier. The fundamental question, however, is whether such functionals exist for the given \( \mathcal{X} \)
and \( \mathcal{Y} \), and if the answer is ‘yes’ then how to find them?

It is well known that if (i) \( \mathcal{X} \) and \( \mathcal{Y} \) are disjoint, (ii) the cardinality, \( |\mathcal{X} \cup \mathcal{Y}| \), of \( \mathcal{X} \cup \mathcal{Y} \) does
not exceed \( n + 1 \), and (iii) elements of \( \mathcal{X} \cup \mathcal{Y} \) are in a general position, then they are vertices of a
simplex. Hence, in this setting, there is always a linear functional \( l \) separating \( \mathcal{X} \) and \( \mathcal{Y} \).
Rosenblatt’s $\alpha$-perceptron [40] used a population of linear threshold elements with random synaptic weights ($A$-elements) as the layer before an $R$-element, that is, a linear threshold element which learns iteratively (authors of some papers and books called the $R$-elements ‘perceptrons’ and lose the complex structure of $\alpha$-perceptron with a layer of random $A$-elements). The randomly initiated elements of the first layer can undergo selection of the most relevant elements.

According to Rosenblatt [40], any set of data vectors becomes linear separable after transformation by the layer of $A$-elements, if the number of these randomly chosen elements is sufficiently large. Therefore, the perceptron can solve any classification problem, where classes are defined by pointing out examples (ostensive definition). But this ‘sufficiently large’ number of random elements depends on the problem and may be large, indeed. It can grow for a classification task proportionally to the number of the examples. The perceptron with a sufficiently large number of $A$-elements can approximate binary-valued functions on finite domains with arbitrary accuracy. Recently, the bounds on errors of these approximations are derived [43]. It is proved that, unless the number of network units grows faster than any polynomial of the logarithm of the size of the domain, a good approximation cannot be achieved for almost any uniformly randomly chosen function. The results are obtained by the application of concentration inequalities.

The method of random projections became popular in machine learning after the Johnson–Lindenstrauss lemma [44], which states that relatively large sets of $m$ vectors in a high-dimensional Euclidean space $\mathbb{R}^d$ can be linearly mapped into a space of much lower dimension $n$ with approximate preservation of distances. This mapping can be constructed (with high probability) as a projection on $n$ random basis vectors with rescaling of the projection with a factor $\sqrt{d}$ [45]. Repeating the projection $O(m)$ times and selecting the best of them, one can achieve the appropriate accuracy of the distance preservation. The number of points $m$ can be exponentially large with $n$ ($m \leq \exp(cn)$).

Two unit random vectors in high dimension are almost orthogonal with high probability. This is a simple manifestation of the so-called waist concentration [13]. A high-dimensional sphere is concentrated near its equator. This is obvious: just project a sphere onto a hyperplane and use the concentration argument for a ball on the hyperplane (with a simple trigonometric factor). This seems highly non-trivial, if we ask: near which equator? The answer is: near each equator. This answer is obvious because of rotational symmetry, but it seems to be counterintuitive.

We call vectors $x$, $y$ from Euclidean space $\mathbb{R}^n$ $\varepsilon$-orthogonal if $|\langle x, y \rangle| < \varepsilon$ ($\varepsilon > 0$). Let $x$ and $y$ be i.i.d. random vectors distributed uniformly (rotationally invariant) on the unit sphere in Euclidean space $\mathbb{R}^n$. Then the distribution of their inner product satisfies the inequality (see, for example, [16] or [46] and compare with the Maxwellian and Lévy’s lemma):

$$P(|\langle x, y \rangle| < \varepsilon) \geq 1 - 2 \exp(-\frac{1}{2} n \varepsilon^2).$$

**Proposition 2.2.** Let $x_1, \ldots, x_N$ be i.i.d. random vectors distributed uniformly (rotationally invariant) on the unit sphere in Euclidean space $\mathbb{R}^n$. For

$$N < \varepsilon^2 n^4 \left[ \ln \left( \frac{1}{1 - \vartheta} \right) \right]^{1/2} \tag{2.1}$$

all vectors $x_1, \ldots, x_N$ are pairwise $\varepsilon$-orthogonal with probability $P > 1 - \vartheta$ [46].

There are two consequences of this statement: (i) in high dimension there exist exponentially many pairwise almost orthogonal vectors in $\mathbb{R}^n$ and (ii) $N$ random vectors are $\varepsilon$-orthogonal with high probability $P > 1 - \vartheta$ even for exponentially large $N$ (2.1). Existence of exponentially large $\varepsilon$-orthogonal systems in high-dimensional spaces was discovered in 1993 by Kainen & Kůrková [47]. They introduced the notion of quasiorthogonal dimension, which was immediately used in the problem of random indexing of high-dimensional data [21]. The fact that an exponentially large random set consists of pairwise $\varepsilon$-orthogonal vectors with high probability was demonstrated in [46] and used for analysis of the data approximation problem in random
bases. We show that not only do such \( \varepsilon \)-orthogonal sets exist, but also that they are typical in some sense.

\( N \) randomly generated vectors \( x_i \) will be almost orthogonal to a given data vector \( y \) (the angle between \( x \) and \( y \) will be close to \( \pi/2 \) with probability close to 1). Therefore, the coefficients in the approximation of \( y \) by a linear combination of \( x_i \) could be arbitrarily large and the approximation problem will be ill-conditioned, with high probability. The following alternative is proved for approximation by random bases.

— Approximation of a high-dimensional data vector by linear combinations of randomly and independently chosen vectors requires (with high probability) generation of exponentially large ‘bases’, if we would like to use bounded coefficients in linear combinations.

— If arbitrarily large coefficients are allowed, then the number of randomly generated elements that are sufficient for approximation is even less than dimension. We have to pay for such a reduction in the number of elements by ill-conditioning of the approximation problem.

We have to choose between a well-conditioned approximation problem in exponentially large random bases and an ill-conditional problem in relatively small (moderate) random bases. This dichotomy is fundamental, and it is a direct consequence of the waist concentration phenomenon. In what follows, we will formally present another concentration phenomenon, stochastic separation theorems \([48,49]\), and outline their immediate applications in artificial intelligence (AI) and neuroscience.

3. Stochastic separation theorems and their applications in artificial intelligence systems

(a) Stochastic separation theorems

Existence of a linear functional that separates two finite sets \( \mathcal{X}, \mathcal{Y} \subset \mathbb{R}^n \) is no longer obvious when \( |\mathcal{X} \cup \mathcal{Y}| \gg n \). A possible way to answer both questions could be to cast the problem as a constrained optimization problem within the framework of, for example, support vector machines \([26]\). The issue with this approach is that theoretical worst-case estimates of computational complexity for determining such functions are of the order \( O(|\mathcal{X} \cup \mathcal{Y}|^3) \) (for quadratic loss functions); a posteriori analysis of experiments on practical use cases, however, suggests that the complexity could be much smaller than \( O(|\mathcal{X} \cup \mathcal{Y}|^3) \) and becomes linear or even sublinear in \( |\mathcal{X} \cup \mathcal{Y}| \) \([50]\).

This apparent discrepancy between the worst-case estimates and a posteriori evaluation of computational complexities can be resolved if concentration effects are taken into account. If the dimension \( n \) of the underlying topological vector space is large, then random finite but exponentially large in \( n \) samples are linearly separable, with high probability, for a range of practically relevant classes of distributions. Moreover, we show that the corresponding separating functionals can be derived using Fisher linear discriminants \([39]\). Computational complexity of the latter is linear in \( |\mathcal{X} \cup \mathcal{Y}| \). It can be made sublinear too if proper sampling is used to estimate corresponding covariance matrices. As we have shown in \([49]\), the results hold for i.i.d. random points from equidistributions in a ball, a cube, and from distributions that are products of measures with bounded support. The conclusions are based on stochastic separation theorems for which the statements for relevant classes of distributions are provided below.
Theorem 3.1 (equidistribution in $\mathbb{B}_n(1)$ [48,49]). Let $\{x_1, \ldots, x_M\}$ be a set of $M$ i.i.d. random points from the equidistribution in the unit ball $\mathbb{B}_n(1)$. Let $0 < r < 1$, and $\rho = \sqrt{1 - r^2}$. Then

$$P\left(\|x\| > r \text{ and } x_i, \frac{x_M}{\|x_M\|} < r \text{ for all } i \neq M\right) \geq 1 - r^n - 0.5(M - 1)\rho^n, \quad (3.1)$$

$$P\left(\|x\| > r \text{ and } x_i, \frac{x_j}{\|x_j\|} < r \text{ for all } i, j, i \neq j\right) \geq 1 - Mr^n - 0.5M(M - 1)\rho^n \quad (3.2)$$

and

$$P\left(\|x\| > r \text{ and } x_i, \frac{x_j}{\|x_j\|} < r \text{ for all } i, j, i \neq j\right) \geq 1 - Mr^n - M(M - 1)\rho^n. \quad (3.3)$$

The proof of the theorem can be illustrated with figure 1. The probability that a single element, $x_M$, belongs to the difference $\mathbb{B}_n(1) \setminus \mathbb{B}_n(r)$ of two $n$-balls centred at $O$ is not smaller than $1 - r^n$. Consider the hyperplane

$$l(x) = r, \text{ where } l(x) = \left(x_i, \frac{x_M}{\|x_M\|}\right).$$

This hyperplane partitions the unit ball $\mathbb{B}_n(1)$ centred at $O$ into two disjoint subsets: the spherical cap (shown as a grey shaded area in figure 1) and the rest of the ball. The element $x_M$ is in the shaded area and is on the line containing the vector $OO'$. The volume of this spherical cap does not exceed the volume of the half-ball of radius $r$ centred at $O'$ (the ball $\mathbb{B}_n(r)$ is shown as a dashed circle centred at $O'$ in the figure). Recall that

$$P(A_1 \& A_2 \& \ldots \& A_m) \geq 1 - \sum_{i} (1 - P(A_i)) \text{ for any events } A_1, \ldots, A_m. \quad (3.4)$$

This ensures that (3.1) holds. Applying the same argument to all elements of the set $S$ results in (3.2). Finally, to show that (3.3) holds, observe that the length of the segment $OA$ on the tangent line to the sphere $S_{n-1}(1)$ centred at $O'$ is always smaller than $r = |OO'|$. Hence the cosine of the angle between an element from $(\mathbb{B}_n(1) \setminus \mathbb{B}_n(r)) \setminus \mathbb{B}_n(r)$ and the vector $OO'$ is bounded from above by $\cos(\angle(OA', OO')) = r$. The estimate now follows from (3.4).

According to theorem 3.1, the probability that a single element $x_M$ from the sample $S = \{x_1, \ldots, x_M\}$ is linearly separated from the set $S \setminus \{x_M\}$ by the hyperplane $l(x) = r$ is at least

$$1 - r^n - 0.5(M - 1)(1 - r^2)^{n/2}.$$

This probability estimate depends on both $M = |S|$ and dimensionality $n$. An interesting consequence of the theorem is that if one picks a probability value, say $1 - \vartheta$, then the maximal
possible values of $M$ for which the set $S$ remains linearly separable with probability that is no less than $1 - \delta$ grows at least exponentially with $n$. In particular, the following holds.

**Corollary 3.2.** Let \( \{x_1, \ldots, x_M\} \) be a set of $M$ i.i.d. random points from the equidistribution in the unit ball \( \mathbb{B}_n(1) \). Let $0 < r, \vartheta < 1$, and $\rho = \sqrt{1 - r^2}$. If
\[
M < \frac{2(\vartheta - r^n)}{\rho^n},
\]
then \( \mathbb{P}(x_i, x_M) < r\|x_M\| \) for all $i = 1, \ldots, M - 1 > 1 - \vartheta$. If
\[
M < \left( \frac{r}{\rho} \right)^n \left( -1 + \sqrt{1 + \frac{2\vartheta \rho^n}{n^2}} \right),
\]
then \( \mathbb{P}(x_i, x_j) < r\|x_i\| \) for all $i, j = 1, \ldots, M, i \neq j \geq 1 - \vartheta$.

In particular, if inequality (3.6) holds, then the set \( \{x_1, \ldots, x_M\} \) is linearly separable with probability $p > 1 - \vartheta$.

The linear separability property of finite but exponentially large samples of random i.i.d. elements is not restricted to equidistributions in \( \mathbb{B}_n(1) \). As has been noted in [22], it holds for equidistributions in ellipsoids as well as for the Gaussian distributions. Moreover, it can be generalized to product distributions in a unit cube. Consider, for example, the case when coordinates of the vectors $x = (X_1, \ldots, X_n)$ in the set $S$ are independent random variables $X_i, i = 1, \ldots, n$ with expectations $\bar{X}_i$ and variances $\sigma_i^2 > \sigma_0^2 > 0$. Let $0 \leq X_i < 1$ for all $i = 1, \ldots, n$. The following analogue of theorem 3.1 can now be stated.

**Theorem 3.3 (product distribution in a cube [49]).** Let \( \{x_1, \ldots, x_M\} \) be i.i.d. random points from the product distribution in a unit cube. Let
\[
R_0^2 = \sum_i \sigma_i^2 \geq n \sigma_0^2
\]
and $0 < \delta < \frac{1}{2}$. Then
\[
\mathbb{P} \left( 1 - \delta \leq \frac{\|x_j - \bar{x}\|^2}{R_0^2} \leq 1 + \delta \text{ and } \left( \frac{x_j - \bar{x}}{R_0}, \frac{x_M - \bar{x}}{\|x_M - \bar{x}\|} \right) < \sqrt{1 - \delta} \text{ for all } i, j, i \neq M \right)
\geq 1 - 2M \exp \left( -\frac{2\delta^2 R_0^4}{n} \right) - (M - 1) \exp \left( -2R_0^4(2 - 3\delta)^2 \right),
\]
and
\[
\mathbb{P} \left( 1 - \delta \leq \frac{\|x_j - \bar{x}\|^2}{R_0^2} \leq 1 + \delta \text{ and } \left( \frac{x_j - \bar{x}}{R_0}, \frac{x_j - \bar{x}}{\|x_j - \bar{x}\|} \right) < \sqrt{1 - \delta} \text{ for all } i, j, i \neq j \right)
\geq 1 - 2M \exp \left( -\frac{2\delta^2 R_0^4}{n} \right) - M(M - 1) \exp \left( -2R_0^4(2 - 3\delta)^2 \right). \tag{3.8}
\]

The proof is based on concentration inequalities in product spaces [14,51]. Numerous generalizations of theorems 3.1 and 3.3 are possible for different classes of distributions, for example for weakly dependent variables, etc.

Linear separability, as an inherent property of datasets in high dimension, is not necessarily confined to cases whereby a linear functional separates a single element of a set from the rest. Theorems 3.1 and 3.3 can be generalized to the case of separating $m$-tuples, $m > 1$ too. An example of such generalization is provided in the next theorem.

**Theorem 3.4 (separation of $m$-tuples [52]).** Let $X = \{x_1, \ldots, x_M\}$ and $Y = \{x_{M+1}, \ldots, x_{M+k}\}$ be i.i.d. samples from the equidistribution in $\mathbb{B}_n(1)$. Let $Y_k = \{x_{M+r_1}, \ldots, x_{M+r_m}\}$ be a subset of $m$ elements
from $\mathcal{Y}$ such that
\[
\beta_2(m - 1) \leq \sum_{i \neq i_r} (x_{M+r_i}, x_{M+r_i}) \leq \beta_1(m - 1) \quad \text{for all } i = 1, \ldots, m.
\] (3.9)

Then
\[
P(\exists \text{ a linear functional separating } \mathcal{X} \text{ and } \mathcal{Y}_c) \geq \max_{\varepsilon \in (0,1)} (1 - (1 - \varepsilon)^n)^{\frac{1}{m}} \left(1 - \frac{\Delta(\varepsilon, m)^2}{2}\right)^M,
\] (3.10)

where
\[
\Delta(\varepsilon, m) = 1 - \frac{1}{m} \left(\frac{(1 - \varepsilon)^2 + \beta_2(m - 1)}{\sqrt{1 + (m - 1)\beta_1}}\right)^2,
\]
subject to:
\[
(1 - \varepsilon)^2 + \beta_2(m - 1) > 0, \quad 1 + (m - 1)\beta_1 > 0.
\]

The separating linear functional is again the inner product, and the separating hyperplane can be taken in the form [52]
\[
l(x) = r, \quad \text{where } l(x) = \left(x, \frac{\bar{y}}{\|\bar{y}\|}\right), \quad r = \frac{1}{\sqrt{m}} \left(\frac{(1 - \varepsilon)^2 + \beta_2(m - 1)}{\sqrt{1 + (m - 1)\beta_1}}\right),
\] (3.11)
and $\varepsilon$ is the maximizer of the nonlinear program on the right-hand side of (3.10), and $\bar{y} = (1/m) \sum_{i=1}^m x_{M+r_i}$. To see this, observe that $\|x_{M+r_i}\| \geq 1 - \varepsilon$, $\varepsilon \in (0,1)$, for all $i = 1, \ldots, m$, with probability $(1 - (1 - \varepsilon)^n)^m$. With this probability the following estimate holds:
\[
\left(\frac{\bar{y}}{\|\bar{y}\|}, x_{M+r_i}\right) \geq \frac{1}{m\|\bar{y}\|}((1 - \varepsilon)^2 + \beta_2(m - 1)).
\]
Hence
\[
\frac{1}{m}(1 + (m - 1)\beta_1) \geq (\bar{y}, \bar{y}) \geq \frac{1}{m}((1 - \varepsilon)^2 + \beta_2(m - 1))
\]
and $l(x)$ in (3.11) is the required functional (figure 1).

If the elements of $\mathcal{Y}_c$ are uncorrelated, i.e. the values of $\beta_1(m - 1), \beta_2(m - 1)$ are small, then the distance from the spherical cap induced by linear functional (3.11) to the centre of the ball decreases as $O(1/\sqrt{m})$. This means that the lower-bound probability estimate in (3.10) is expected to decrease too. On the other hand, if the elements of $\mathcal{Y}_c$ are all positively correlated, i.e. $1 \geq \beta_1 > \beta_2 > 0$, then one can derive a lower-bound probability estimate which does not depend on $m$.

The peculiar properties of data in high dimension, expressed in terms of linear separability, have several consequences and applications in the realm of AI and machine learning; examples are provided in the next sections.

(b) Correction of legacy artificial intelligence systems

Legacy AI systems, i.e. AI systems that have been deployed and are currently in operation, are becoming more and more widespread. Well-known commercial examples are provided by global multi-nationals, including Google, IBM, Amazon, Microsoft and Apple. Numerous open-source legacy AIs have been created to date, together with dedicated software for their creation (e.g. Caffe [53], MXNet [54], Deeplearning4j [55] and Tensorflow [56] packages). These AI systems require significant computational and human resources to build. Regardless of resources spent, virtually any AI and/or machine learning-based systems are likely to make a mistake. Real-time correction of these mistakes by retraining is not always viable due to the resources involved. AI retraining is not necessarily desirable either, because AI’s performance after retraining may not always be guaranteed to exceed that of the previous performance. We can, therefore, formulate the technical requirements for the correction procedures. The corrector should: (i) be simple; (ii) not change the skills of the legacy system; (iii) allow fast non-iterative learning; and (iv) allow correction of new mistakes without destroying previous corrections.
A possible remedy to this issue is the AI correction method [22] based on stochastic separation theorems. Suppose that at a time instance $t$ values of signals from inputs, outputs and the internal state of a legacy AI system could be combined together to form a single measurement object, $x = (x_1, \ldots, x_n)$. All $n$ entries in this object are numerical values, and each measurement $x$ corresponds to a relevant decision of the AI system at time $t$. Over the course of the system’s existence a set $\mathcal{S}$ of such measurements is collected. For each element in the set $\mathcal{S}$ a label ‘correct’ or ‘incorrect’ is assigned, depending on external evaluation of the system’s performance. Elements corresponding to ‘incorrect’ labels are then filtered out and dealt with separately by an additional subsystem, a corrector. The process is illustrated in figure 2. In this diagram, the original legacy AI system (shown as legacy AI system 1) is supplied with a corrector altering its responses. The combined new AI system can in turn be augmented by another corrector, leading to a cascade of AI correctors (figure 2).

If distributions modelling elements of the set $\mathcal{S}$ are, for example, an equidistribution in a ball or an ellipsoid, a product of measures distribution, a Gaussian, etc. then

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- Theorems 3.1–3.4 guarantee that construction of such AI correctors can be achieved using mere linear functionals.
- These linear functionals admit closed-form formulae (Fisher linear discriminant) and can be determined in a non-iterative way.
- The availability of explicit closed-form formulae in the form of a Fisher discriminant offers major computational benefits as it eliminates the need to employ iterative and more computationally expensive alternatives, such as support vector machines.
- If a cascade of correctors is employed, the performance of the corrected system drastically improves [22].

The results, perhaps, can be generalized to other classes of distributions that are regular enough to enjoy the stochastic separability property.

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Figure 2. Cascade of AI correctors. (Online version in colour.)
The corrector principle has been demonstrated in [22] for a legacy AI system in the form of a convolutional neural network trained to detect pedestrians in images. AI errors were set to be false positives, and the corrector system had to remove labelled false positives by a single linear functional. A detailed description of the experiment is provided in [22], and a performance snapshot is shown in figure 3. Dimensionality $n$ of the vectors $x$ was 2000. As we can see from figure 3, single linear functionals are capable of removing several errors of a legacy AI without compromising the system’s performance. Note that AI errors, i.e. false positives, were chosen at random and have not been grouped or clustered to take advantage of positive correlation. (The definition of clusters could vary [27].) As the number of errors to be removed grows, performance starts to deteriorate. This is in agreement with our theoretical predictions (theorem 3.4).

(c) Knowledge transfer between artificial intelligence systems

Legacy AI correctors can be generalized to a computational framework for automated AI knowledge transfer whereby labelling of the set $\mathcal{S}$ is provided by an external AI system. AI knowledge transfer has been the focus of growing attention during the last decade [57]. Application of stochastic separation theorems to AI knowledge transfer was proposed in [52], and the corresponding functional diagram of this automated set-up is shown in figure 4. In this set-up a student AI, denoted as $\text{AI}_s$, is monitored by a teacher AI, denoted as $\text{AI}_t$. Over a period of activity system $\text{AI}_s$ generates a set $\mathcal{S}$ of objects $x$, $x \in \mathbb{R}^n$. The exact composition of the set $\mathcal{S}$ depends on the task at hand. If the outputs from $\text{AI}_s$ differ from those of $\text{AI}_t$ for the same input, then an error is registered in the system. Objects $x \in \mathcal{S}$ associated with errors are combined into the set $\mathcal{Y}$. The process gives rise to two disjoint sets:

$$\mathcal{X} = \{x_1, \ldots, x_M\}, \quad \mathcal{X} = \mathcal{S} \setminus \mathcal{Y} \quad \text{and} \quad \mathcal{Y} = \{x_{M+1}, \ldots, x_{M+k}\}.$$  

Having created these two sets, knowledge transfer from $\text{AI}_t$ to $\text{AI}_s$ can now be organized in accordance with algorithm 1. Note that data regularization and whitening are included in the preprocessing step of algorithm 1. The algorithm can be used for AI correctors too. Similar to AI correction, AI knowledge transfer also be cascaded. Specific examples and illustrations of AI knowledge transfer based on stochastic separation theorems are discussed in [52].
Stochastic separation theorems are a generic phenomenon, and their applications are not limited to AI and machine learning systems. An interesting consequence of these theorems for neuroscience has been discovered and presented in [58]. Recently, it has been shown that in humans new memories can be learnt very rapidly by supposedly individual neurons from a limited number of experiences [59]. Moreover, neurons can exhibit remarkable selectivity to complex stimuli, the evidence that has led to debates around the existence of the so-called ‘grandmother’ and ‘concept’ cells [60–62], and their role as elements of a declarative memory. These findings suggest that not only can the brain learn rapidly but also it can respond selectively to ‘rare’ individual stimuli. Moreover, experimental evidence indicates that such a cognitive functionality can be delivered by single neurons [59–61]. The fundamental questions, hence, are: How is this possible? and What could be the underlying functional mechanisms?

It has been shown in [58] that stochastic separation theorems offer a simple answer to these fundamental questions. In particular, extreme neuronal selectivity and rapid learning can already be explained by these theorems. Model-wise, the explanation of extreme selectivity is based on a conventional and widely accepted phenomenological generic description of neural response to stimulation. Rapid acquisition of selective responses to multiple stimuli by single neurons is ensured by classical Hebbian synaptic plasticity [63].

4. Conclusion

Twenty-three Hilbert’s problems created important ‘focus points’ for the concentration of efforts of mathematicians for a century. The sixth problem differs significantly from the other twenty-two problems. It is very far from being a purely mathematical problem. It seems to be impossible to imagine its ‘final solution’. The sixth problem is a ‘programmatic call’ [64], and it works:

— We definitely know that the sixth problem had great influence on the formulation of the mathematical foundation of quantum mechanics [3] and on the development of axiomatic quantum field theory [65].
— We have no doubt (but the authors have no direct evidence) that the sixth problem has significantly affected research in the foundation of probability theory [4] and statistical mechanics [5].
— The modern theory of measure concentration phenomena has direct relations to the mathematical foundations of probability and statistical mechanics, uses results of Kolmogorov and Khinchin (among others), and definitely helps to create ‘a rigorous and satisfactory development of the method of the mean values...’ [1, p. 454].
Algorithm 1 AI knowledge transfer/correction [52].

(i) Preprocessing

(a) Centring. For the given set $S$, determine the set average, $\overline{x}(S)$, and generate sets $S_c$

$$S_c = \{ x \in \mathbb{R}^n | x = \xi - \overline{x}(S), \ \xi \in S \},$$

$$\mathcal{Y}_c = \{ x \in \mathbb{R}^n | x = \xi - \overline{x}(S), \ \xi \in \mathcal{Y} \}.$$

(b) Regularization. Determine covariance matrices $\text{Cov}(S_c)$, $\text{Cov}(S_c \setminus \mathcal{Y}_c)$ of the sets $S_c$ and $S_c \setminus \mathcal{Y}_c$. Let $\lambda_i(\text{Cov}(S_c))$, $\lambda_i(\text{Cov}(S_c \setminus \mathcal{Y}_c))$ be their corresponding eigenvalues, and $h_1, \ldots, h_n$ be the eigenvectors of $\text{Cov}(S_c)$. If some of $\lambda_i(\text{Cov}(S_c))$, $\lambda_i(\text{Cov}(S_c \setminus \mathcal{Y}_c))$ are zero or if the ratio $\frac{\max_i |\lambda_i(\text{Cov}(S_c))|}{\min_i |\lambda_i(\text{Cov}(S_c))|}$ is too large, project $S_c$ and $\mathcal{Y}_c$ onto an appropriately chosen set of $m < n$ eigenvectors, $h_{n-m+1}, \ldots, h_n$:

$$S_r = \{ x \in \mathbb{R}^n | x = H^T \xi, \ \xi \in S_c \},$$

$$\mathcal{Y}_r = \{ x \in \mathbb{R}^n | x = H^T \xi, \ \xi \in \mathcal{Y} \},$$

where $H = (h_{n-m+1} \cdots h_n)$ is the matrix consisting of $m$ significant principal components of $S_c$.

(c) Whitening. For the centred and regularized dataset $S_r$, derive its covariance matrix, $\text{Cov}(S_r)$, and generate whitened sets

$$S_w = \{ x \in \mathbb{R}^n | x = \text{Cov}(S_r)^{-\frac{1}{2}} \xi, \ \xi \in S_r \},$$

$$\mathcal{Y}_w = \{ x \in \mathbb{R}^n | x = \text{Cov}(S_r)^{-\frac{1}{2}} \xi, \ \xi \in \mathcal{Y}_r \}.$$

(ii) Knowledge transfer

(a) Clustering. Pick $p \geq 1$, $p \leq k$, $p \in \mathbb{N}$, and partition the set $\mathcal{Y}_w$ into $p$ clusters $\mathcal{Y}_{w,1}, \ldots, \mathcal{Y}_{w,p}$ so that elements of these clusters are, on average, pairwise positively correlated. That is, there are $\beta_1 \geq \beta_2 > 0$ such that:

$$\beta_2(|\mathcal{Y}_{w,i}| - 1) \leq \sum_{\xi \in \mathcal{Y}_{w,i}} (\xi, x) \leq \beta_1(|\mathcal{Y}_{w,i}| - 1) \text{ for any } x \in \mathcal{Y}_{w,i}.$$

(b) Construction of auxiliary knowledge units. For each cluster $\mathcal{Y}_{w,i}$, $i = 1, \ldots, p$, construct separating linear functionals $\ell_i$ and thresholds $c_i$:

$$\ell_i(x) = \left( \frac{w_i}{\|w_i\|}, x \right),$$

$$w_i = \left( \text{Cov}(S_w \setminus \mathcal{Y}_{w,i}) + \text{Cov}(\mathcal{Y}_{w,i}) \right)^{-1} (\overline{x}(\mathcal{Y}_{w,i}) - \overline{x}(S_w \setminus \mathcal{Y}_{w,i})), $$

$$c_i = \min_{\xi \in \mathcal{Y}_{w,i}} \left( \frac{w_i}{\|w_i\|}, \xi \right),$$

where $\overline{x}(\mathcal{Y}_{w,i})$, $\overline{x}(S_w \setminus \mathcal{Y}_{w,i})$ are the averages of $\mathcal{Y}_{w,i}$ and $S_w \setminus \mathcal{Y}_{w,i}$, respectively. The separating hyperplane is $\ell_i(x) = c_i$.

(c) Integration. Integrate auxiliary knowledge units into decision-making pathways of AIx. If, for an $x$ generated by an input to AIx, any of $\ell_i(x) \geq c_i$ then report $x$ accordingly (swap labels, report as an error, etc.)

— The modern idea of blessing of dimensionality in high-dimensional data analysis [18,19,48] is, in essence, an extension and further development of ideas from the mathematical foundations of statistical mechanics.
The classical measure concentration theorems state that random points in a highly dimensional data distribution are concentrated in a thin layer near an average or median-level set of a Lipschitz function. The stochastic separation theorems describe the fine structure of these thin layers: the random points are all linearly separable from the rest of the set even for exponentially large random sets. Of course, for all these concentration and separation theorems the probability distribution should be ‘genuinely’ high dimensional. Equidistributions in balls or ellipsoids or the products of distributions with compact support and non-vanishing variance are simple examples of such distributions. Various generalizations are possible.

For which dimensions does the blessing of dimensionality work? This is a crucial question. The naive point of view that the dimension of data is just a number of coordinates is wrong. This is the dimension of the data space, where data are originally situated. The notion of intrinsic dimension of data is needed [66,67]. The situation when the number of data points \( N \) is less (or even much less) than the dimension \( d \) of the data space is not new. Moreover, Donoho [18] considered the property \( d > N \) as a generic case in the ‘post-classical world’ of data analysis. In such a situation, we really explore data on a \( d - 1 \)-dimensional plane and should modestly reduce our high-dimensional claim. The projection of data on that plane can be performed by various methods. We can use as new coordinates projections of points on the known data points or Pearson’s correlation coefficients, when they are suitable; for example, when the data points are fragments of time series or large spectral images, etc. In these new coordinates the data table becomes a square matrix and further dimensionality reduction could be performed using principal component analysis (PCA), or its nonlinear versions like principal manifolds [28] or neural auto-encoders [68].

A standard example can be found in [69]: the initial data space consisted of fluorescence diagrams and had dimension \( 5.2 \times 10^5 \). There were 62 data points, and a combination of correlation coordinates with PCA showed intrinsic dimension 4 or 5. For the selection of relevant principal components the Kaiser rule, the broken stick models or other heuristical or statistical methods can be used [70].

A similar preprocessing ritual is helpful even in more ‘classical’ cases when \( d < N \). The correlation (or projection) transformation is not essential here, but the formation of relevant features with dimension reduction is important. If after model reduction and whitening (transformation of coordinates to get the unit covariance matrix, step 1.c in algorithm 1) the new dimension \( D \geq 100 \), then for \( \leq 10^6 \) data points we can expect that the stochastic separation theorems work with probability greater than 99%. Thus the separation of errors with Fisher’s linear discriminant is possible, and many other ‘blessing of dimensionality benefits’ are achievable. Of course, some additional hypotheses about the distribution functions are needed for a rigorous proof, but there is practically no chance to check them \textit{a priori} and the validation of the whole system \textit{a posteriori} is necessary. In smaller dimensions (for example, less than 10), nonlinear data approximation methods can work well, capturing the intrinsic complexity of data, like principal graphs do [29,71].

We have an alternative: either essentially high-dimensional data with thin shell concentrations, stochastic separation theorems and efficient linear methods, or essentially low-dimensional data with efficient complex nonlinear methods. There is a problem of the ‘no man’s land’ in between. To explore this land, we can extract the most interesting low-dimensional structure and then consider the residual as an essentially high-dimensional random set, which obeys stochastic separation theorems. We do not currently know a theoretically justified efficient approach to this area, but here we should say, following Hilbert: ‘Wir müssen wissen, wir werden wissen’ (We must know, we shall know).

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