

perhaps Late 1951 or early 1952 - Relatively forsterian!

T. M. Review (T.M.R.)

This will be a summary-explanation of the history of the analysis, starting from basic concepts. The missions and the reasons for various investigations and "study problems" will be given.

Outline of important ideas:

- 1) What is probability? Manyman
- 2) " " Life?
- 3) How does evolution proceed?
- 4) What is "thinking"? How can one judge when a machine is "thinking"? Give some intuitive ideas on this. Give a few problems that a T.M. should be able to solve: e.g., synthetic geometry problems. Also problems in some general field should be solved more quickly after machine has solved n problems. This should all be done in an un-ad-hoc-way. I shall be the judge of ad-hocness.
- 5) Difference between living being and thinking-machine, i.e. the goal of life is fertile reproduction; the goal of T.M. is quick solution of the problems fast to it.
- 6) How does memory work? Just what determines what it is about a particular situation that we remember?
- 7) Learning and conditioning. Just what determines how a present experience will modify future behavior? This is linked to the problem of memory.
- 8) Antibodies. How do antibodies adapt themselves to destroy a particular kind of germ? How does previous "experience" with that germ help? This is a simple, low life level case of adaptation and learning.
- 9) Adaptation: How do organisms learn to adapt to complex environments in ways to further their goals?
- 10) Languages - how do they grow, and how does one determine what the words will be.
- 11) How does one make a decision from several conflicting lines of evidence?

Started
in 1951, or early 1952,
I believe.

Ap. 2, 1952

①
2

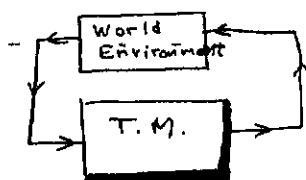
Up to ≈ 22 by middle of 1952 (20 is Feb 6, 1952) ≈ 16 is March 1, 1952
 ≈ 22 Ap. 2, 1952

T. M. — A Survey — *

The simplification of the T. M. problem lies in its parallel statement in various fields of inquiry. Each of these fields has an apparently different problem, yet any progress in one of these problems is translatable into progress in each of the others. It will be the purpose of this paper to outline each of the problems in the various fields, and show how the parts of the respective problems correspond to one another.

① The first problem is the Thinking machine proper. To construct, with existing equipment, a machine that will "think".

More exactly:



— T. M. Has an input and an output — These may be of any form that conveys information.

— T. M.'s output goes into the input of the "world Environment", and the W.E.'s output is the input of T. M..

— Now T. M. has a certain set of inputs which it regards as "good" — with varying degrees of goodness.

— The purpose of T. M. is to give an output such that ~~deserves~~ the total "goodness" of its input is maximized. To do this, it must make experimental outputs and observations on its resultant input — in order to determine the nature of W.E.

The exact nature of T. M.'s "goodness" criterion is rather uncertain. Suppose it associates with each input $I(t)$ at time t , an amt. of good, $G(I(t))$,

— and T. M. tries to maximize some $\int_0^t F(t, G(I(t))) dt$

— where F is some func. of G and t , and the integration limits are birth and death of T. M. [Note: $\frac{d}{dt}$ and $\frac{d^2G}{dt^2}$ maybe under the F .]

— What T. M. must do at each moment, is to act so as to maximize the expected value of this F !

— This involves a definition of probability. T. M. must be able to associate with every future plan of action,

— answer questions and very possible value of the S_i ,

— & probability that density, that such a course of action will ~~lead~~ to such a value of the S_i .

T.M.

— But this is an excessive simplification, since one will
make decisions in the future, which cannot will be made
 — on the basis of information not now available.

What one does is associate each "plan of behavior,"
 — ~~an~~ "integral value," pair, with a probability.

— A "plan of behavior" is a system ~~designed~~ which
 tells one what to do at time t , upon the
 basis of all inputs ~~up~~ ~~to~~ time t . [This concept
 is identical to one used ~~in~~ in the "Theory of Games"]

E.G. Given a "plan of behavior" for scrutiny, T.M. must
 be able to tell the probability that the "good" integral
 will be between .7 and .8.

But, may these "behavior plans" include such statements as
 "put the following plans into the plan evaluator!": 13, 12 and 27,
~~and~~ start at time 178, carry out the ~~one~~ with the
 highest expected ~~value~~ value of the "Good's"?

This problem seems to be identical with the one about
 "expectancy maximization in a Capitalistic Economy."

(2) The problem of "what is probability?" Given a ~~series~~
 set of objects. Given a series of them. What is the probability
 that the $n+1^{\text{th}}$ is a certain one of the set?

Actually, probability is an elementalistic concept. Prob. is
 used to divide up a more complex problem into solvable parts.

In some problems where elementalistic ~~calculus~~ methods fail,
 one occasionally can ask for a probability where no such
 probability exists.

Also, there seems to be the idea of "certainty of probability"
 or "probability of probability". Since probability is
 gotten by observing a series of events, one would expect
 a short series to yield a less certain value of ~~prob.~~
 "mean freq." or "probability" than a long series.

The crucial factor in prob. is abstraction. One
 observes the first n events. One observes the frequency
 of abstractions A, B, and C, upon those events. Now,

say element ϕ_i has ~~the~~ abstractions A, B and C.
 To find the prob. of event ϕ_i at the $n+1^{\text{th}}$ time

period, one uses the ~~the~~ observed frequencies
 of the abstractions A, B and C. not given these freqs, what is the prob. that they will occur together is an unsolved problem.

See footnote
 at bottom
 of page

3

(3) The problem "How may abstractions be most readily and naturally be constructed?" This is a key to the problem of probability as stated above, tho it is not the whole solution. That generalized concept of abstraction is simply a listing of all the objects in the universe that have that abstraction in common.

Perhaps the important problem is, given an event, which abstractions from this event will most probably be most useful? — Perhaps a soln. like to this is afforded by the following: Consider the set of all abstraction methods — now non-El. those of the set that give highest reliability in probability estimates — i.e. those which are most significant in signaling out events or, less elementarily, those abstraction methods which have, in the past "led" to greatest increase of the "good S", are the best methods to use. ("in the past" leads to ad-hoc methods).

(4) The Problem of Life: Each living organism is a device which must reproduce, in order to survive as a species.

Reproduction, then, is the ultimate "good" for a living I. M., and the number of fertile offsprings is a measure of exactly the "good". Actually, the exact problem of "good" for a living creature, is a difficult one. Essentially, it implies the ~~max~~ largest number of fertile descendants, in the "long run" — but statement of this criterion in terms of the individual T.M., is more difficult. Maximum no. of fertile offsprings is a good approx. (It clearly, a fertile offspring with sterile offsprings is ~~useless~~ which explains why this is only a rough approx.)

^{see also bottom of page 7} Footnote to top of page (2). The problem is: shall one consider action based on expectancy of a) expected causes of action or b) expected "plans of action" or c) "expected" plans of action" which include reference to the plan evaluator or d) some more complex method? Is method a) actually the most general possible — — presupposing that re-evaluation is made at each step in time? Very probably, the following is a soln: (to page 4)

- Q - The problems of learning and memory: These are spoken of together, since there is probably a close connection.

In memory, we have input data from the world. This data

- is abstracted before it is stored (.probably). - Data must be filed in such a way that it will be useful in future problem solving. The manner of filing (what abstractions are used for filing) is quite important, and what factors determine the manner in which an item is to be stored, - is important.

Also, in conditioned response "learning", we have abstractions made - what are these abstractions? - Also what governs

Learning rate?

^E footnote, continued: Consider T.M. and The Environment.

from
Bottom
of (3)

"having discrete, alternate moves, as in chess. - Let an "Action plan" be ~~weakly~~^{for T.M.} defined by the following: A ~~for each~~^{correspondence betw.} set of moves by Env. and a corresp. set by T.M. in reply.

~~E~~. These correspondences are limited in that - if 2 sets - - of Env... moves are identical up to time to_r, then T.M.'s theme - polyph. in the "Action plan" must be identical up to

-- An "action-plan" for Env. may be similarly defined -

— Each action plan has a certain time after which it is to come into effect.

- An "action plane" for T.M. and one for E.n.
both defined after t_2 , define a game after
 t_2 .

-- Now say the "good integral" is a functional on
the set of games.

Let $A_{T.M.}^i$ be the set of ~~the~~ T.M.s "action plans".

- Post - After time t . Let A_E^{t+j} be the Env's " " ...
 " " " " $t+1$. Let $A_{\text{sys}}^{t+j} A_{\text{TM}}^{t+j}$ be the game after
 the resulting from these 2 plays.

Let $P(+^t A_e, X^k)$ be the probability (based on observations of the game up to $t-1$) that if T.M. does move X^k at time t then Enr. will use Action-Plan $+^t A_e$.

Let $G(t^+ A_E^2 + A_{T.M}^2)$ be the value of the good functional for the entire game considered - the known game up to $t=1$, and the game defined by $t^+ A_E^2 + A_{T.M}^2$ after time $t=1$.

$$\text{Then } \sum_j P(t+1 A_E^j, X^K) G(t+1 A_E^j, A_{TM})$$

(where Σ runs over all values of j , and j runs over ~~all~~
only those plus which begin with the ~~the~~ move X^K)

is the expectancy in G , that will result from the move X^K .

We want to choose X^K to maximize this expectancy.

Note: I think that this solution is rather non-elementalistic.

After the game starts, an action plan can be derived by T.M. Actually, before the game starts, an action plan can be derived by T.M.

are computed ~~once~~ and ~~for all time~~ no computation need be performed this problem after that time, or at each move. Computation of is quite similar to the chessboard problem.

the probabilities of each of Env's action plans is a very difficult task. It involves at ~~each move~~ of each plan,

playing Machine problems terms of the past. [Note: Actually, before the game starts, T.M. computes the action plan, once and for all time. At this time, T.M. has some memory of shannons.]

A-priori, Env's first move is arbitrary, and probably and can be ~~almost~~ arbitrary. But as the

Env's ~~2nd~~ move is also ~~almost~~ arbitrary. Similarly, Env continues. T.M. begins to see possible systems in Env's moves, and these systems make better probability estimates possible.

Actually, the above soln. seems to be complete — "action plans" describe T.M.'s and Env's actions completely as is possible. It is desirable to elementalize the computation of the probabilities of Env's "action plans".

Footnote to (4), page 3: The problem of "what are the desirable traits that evolution selects for" is the problem of "what is the 'good functional' for life" (as referred to abstractly on page 4)?

Actually, in the case of social animals, not the individual, but the society as a whole, is selected for, by evolution. Here we can treat the society as a whole, as a T.M., with a certain "good functional" (its reproduction rate), which it must maximize — The problem of T.M. then reduces to the following:

Given the moves of the game up to t , what is the probability that Env's move at $t+1$ will be X^K ?

(6) The Problem of "Scientific Method": In science one must make a-priori guesses at theories. Given

- a set of phenomena, one can make an inf. no. of theories that satisfy ~~all~~ all of those phenomena. Out of all of those theories, one must choose the "most probable" ones — i.e. those which will ^{be} most likely to give valid extrapolations of the observed phenomena. Then after choosing the "most probable" theories, one tests them empirically. — This whole problem is identical with the problem of "What is probability?" (see Q. page 2)

⑦ The problem of Antibodies, or Adaptation: — Normally, with inanimate life, adaptation is by conforming. E.g., if a river runs thru a gorge, the gorge will wear away. Contrast this with a tire shoe. Instead of wearing away the skin, a callouse (thicker skin) may form. — Similarly, it seems that anti-body production is a simplified kind of "negative conforming". that is typical of "life". — Instead of the animal giving in to the insurgent organisms, the animal's blood puts up a fight. — Actually this entire problem isn't very clearly stated, and its connection with "life" should be made clear. It is felt that "negative conforming" shows the great and versatile adaptability to environment that is so characteristic of life.

Summary so far: It can be seen now that problems ④ (Thinking Machine) ⑤ (Probability) and ⑥ (Scientific method) are intimately related and identical. ③ (Methods of Abstraction) is related to ⑤ (Prob.) in the following manner: One has a game up to time t_0 to find ~~prob.~~ prob. of Env's move X^k : X^k is a member of abstraction A_x . Now in the past, it has been found that if three successive moves were a member of abstraction A_3 , then the prob. that the next move would be a member of A_{x_3} is .25, with ~~allowing~~ a certain number of observed cases (say 23). Now the t_0-2 , t_0-1 and t_0 moves are members of A_3 ; so the prob. of X^k at t_0 is .25 — from this abstraction alone. Methods of combining data from various abstractions is a difficult problem (see Q. p 48 for) an attempt

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of separation is made
"preferable", if it
is found to lead, soon,
to an increase in "good" — This kind of, but it may be a condition that
is almost always found to be true.

2

→ presupposes a certain kind of
G, that is in the form of an
S, or some similar condition —

Here, again, we have the problem of ad-hoc abstractions.
If we eliminate the possibility of an "origin" in time
(a natural zero point), then this reduces the poss. of
Many ad-hoc hypotheses. Also, for a very long
run of moves (and all "life" has the moves of the ancestors
incorporated into its system to some degree) it becomes
almost impossible to describe certain ad-hoc abstractions
with the limited, finite nervous system of the organism.
Perhaps this idea, with the idea of abstraction/^{methods} on
page 3, will solve the problem. [see bottom of 8]

Note: By taking simple T.M.'s like * coupling's
or the Homostat ^{its} various orders of complexity, one
can, by working backward, define probability.

Also, for any simple definition of probability
one can construct a T.M. — Now then, simplify or
otherwise improve this T.M., then go back to
the prob. it defines, simplify and/or improve that
def. of prob., then go to its T.M. and improve it —
and so on, back and forth. — This is a trick that
can be worked with between any of the many
aspects of this problem.

Try to construct a T.M. or Homostat or a def. of
prob. based on Wren's Affective tone mechanism.

I did devise a T.M. for this mechanism. — by
using rate of change of "Good" as well as ^{absolute} amount
of good, this T.M. can be improved — see what
kind of prob. it leads to.

Note: Chess or Checkers, to show th. redundancy of:
for the first move, White must chose an action
plan. Either 1 there exists an action plan such that
Black always loses or 2 there exists only action plans
in which Black occasionally wins.

If 1 is true, White will, if he is "perfect player",
choose a winning plan. If 2 is true, and Black is
perfect, then Black will win. [this does not prove it; see p. 15.1]
from ④ Page ③. The problem of what characteristics of
an individual are maximized so that net reproduction rate is
maximized is helped along by the following.

For each individual construct, for the nth generation,
an n-dimensional matrix. the $(z_1 z_2 z_3 \dots z_n)$ element -

②

of this matrix will be 1, if the a_n^m offspring of the a_{n-1}^m offspring of the a_{n-2}^m offspring ... of the a_2^m offspring of the a_1^m offspring of the individual exists. That element will be zero if that ~~the~~ offspring does not exist. (order of offsprings is in order of time of conception, since ~~the~~ 2 offsprings of a man by 2 women, may be born simultaneously)

Now consider the aggregate of all such possible matrices. (all such matrices are limited — i.e. if the a_r^m offspring exists, then the a_{r-1}^m offspring must exist). Each matrix characterizes completely the possible character of the n^m generation.

To each such matrix assign a probability, P , that this matrix will represent the true state of the individuals n^m order ancestors

If an animal is limited to A offsprings (A is quite large for most animals if artificial insemination is allowed), the ~~the~~ n^m ancestral matrix has A^n elements. The set of all possible n^m ancestral configurations are of number 2^{A^n} . So the probability distribution will be a 2^{A^n} dimensional vector.

~~all~~ of these vector components will be independent, since all 2 matrices are identical

or we may look upon the probabilities as a scalar field, say, the A^n dimensional vectorspace, where each vector component has the values 0 or 1, only.

[Sectio[n] of 7] Probability in a T.M.: Start out with any "reasonably good" method of prob. evaluation. Then use this method to find a better method of prob. evaluation. And use the "better" method to find a better one yet, etc. Now there is a minimum "reasonable goodness" that is required for this to be possible, also, all initial methods may not converge to the same, or to equally good methods.

This "bootstrap method" operates as follows:

Take the original method, then modify it arbitrarily, and determine the "probability" (by the initial method) that this modification is an improvement. Try different modifications until one is found which is ~~not~~ very probably an improvement, and test it empirically. This may be a very slow method of convergence, but "mother

"nature" has had so much time to work at it that it may be ^{an} O.K. method for her.

(8) The information Packing problem. See ~~Fig.~~ T 29 for discussion. Here, the problem is to construct a language that is as "compact" as possible. Information theory gives / meaning to "^{exact} compactness". The idea of constructing a language is to have abstractions, and methods of combining abstractions (see (3)) — probably in ways other than the Boolean algebra way.

If (3) is solved, we can solve (8) in the following manner. First start out with a basic set of abstractions. Now, whenever ^{not necessarily basic} any abstraction is used (must it be used in what turns out to be an expedient manner?), keep a running account of the freq. of use of each abstraction. We have (by (3)) a method of combining old abstractions to produce new ones — Use this method in a random way, utilizing the most frequently used of the old abstractions. (Note: if sentences (abstractions) are constructed at random, the short (containing few old abstractions) ~~new~~ new abstractions are most likely. Consider that one of the abstractions used is "end abstraction process now" — if the prob. of this abstraction being chosen is p , then the prob. of ~~most~~ a new abstraction ~~also~~ containing n old abstractions is $(1-p)^n p$) The hyper ^{to} a priori preference to short abstractions is in accord with hyper ^{to} a priori probability of ~~other~~ briefly stated scientific laws.

We have a bunch of info. to express via our abstractions, and we try various abstraction methods, to find some (or one) that will most consistently express our data. The ad-hoc abstraction would, of course express our data perfectly, and most briefly, but if we must build our abstractions probabilistically from old abstractions, then our ad-hoc abstraction is extremely improbable.

Feb 6, 1952

This is a return to a survey of T.M.

Thinking about T.M., at first, seems an imposs. task.

It appears that because of Nature's method of developing T.M. being optimal [i.e. to repeat, if ϕ_{ij} is the T.M. op. (ϕ_{ij} is the j^{th} offspring of the i^{th} generation - all T.M.'s reproduce ~~at~~^{on} Jan 1 only each year) - each T.M. dies as soon as it reproduces] the number of offsprings of each T.M. is in the "Goodness" of that partic. operator form. Each set of offsprings is a Gauss distrib. about th. parent in Operator space.

This seems to be a very good way to develop T.M.'s, if one has many resources. - Perhaps it is, in a sense, th. best way!

Even if man had th. resources, he would be limited because he doesn't know th. "metric" of Op. space - i.e. for a gen. Op., how probable is a mutation to another op?

This is th. problem of structure of genes

To repeat: Genetic structure determines the "metric" of Op. space. By knowing it, one could plot a probabilistic distribution of T.M. on, say the N^{th} generation.

To do this - (1) One has th. metric of op. space
(2) one has an environment defined by a transition point function on op. space, giving th. no. of kids for that point in op. space.

~~REMARKABLE~~ Suppose th. op. space could be so distorted that "distance" was ~~very~~ identical with Geom. distance.

For simplicity, let op. space be 1 dimens.

Let $P_n(x, k)$ be th. probability that upon the n^{th} generation, there will be k individuals alive, of operator structure, x . ~~REMARKABLE~~ Then, if one knows ~~what~~ $O(x)$, the no. of ~~at~~ kids x will have in th. gen environment, one can, from $O(x)$ and $P_n(x, k)$, find $P_{n+1}(x, k)$.

Note: Here we assume no interaction between operators after birth. - Certainly, recognition of such a factor would change the soln. entirely.

We have taken th. simplest problem and we have, in principle, written a matrix!

(11)

2

expression for its soln.

Now, can we add any complexities that make th. problem more interesting and/or useful?

→ There is changing environment.

There is simple ~~→~~ Malthusian interaction, in which $p_n(x, k)$ is ~~added~~ multiplied by $\left[\int p_n(x, k) dx dk \right]^{-1}$ for large k to normalize $p_n(x, k)$ - so as to keep th. expected population constant in accord with a limited food supply. Since $p_n(x, k)$ seems to increase exponentially with n , probably this constant renormalization would result in $\rightarrow p_n(x, k)$ consisting of a few S functions in $x = \underline{x_0}$ since th. p_n , resulting from such a distrib. would ~~not~~ $= p_n$. Th. $\underline{x} = 1/b$. condition gives, with renormalization, $p_{\infty} = p_n$.

It appears that Nature had a good metric to start with, and/or a good initial point.

Side Note: A realistic added complexity is $p(x, l)$, the probability that x will have l offsprings in th. g.n. environment.

This has some relevance to th. black box problem, with time delay.

I.E. one puts an input into a black box at time t , and gets an output at time $t+1$. One can put in 10 inputs per second. One tries to maximize the output. (As in Ub 50)

Z 12-1

The changing environment is important.

If th. environment changes too rapidly per generation, th. adaptation criteria for that generation, are not relevant for the next generation, and we lose th. effectiveness of our metric.

A serious problem: for a known metric, and a given starting pt. in sp. space, under what conditions will (a) $\lim_{n \rightarrow \infty} \int p_n(x, k) dx dk \rightarrow \infty$

(b) $\lim_{n \rightarrow \infty} \int p_n(x, k) dx dk \rightarrow 0$?

Just how does a rapidly changing environment make (b) more probable?

This enables us to ask - how good must a metric, init. condns, and environment be, for (a) to occur?

Z 12-7

Z 11-25

from page Z 11, line 25.

The opportunity of having 10 investigations running simultaneously means that when one or 2 investigators have good luck, more man power is allocated to their region of space. This is similar to the no. of kids being of the "goodness" of a pt. in sp. space, [kids = offsprings]

Z 11-50 A good method of analysis is to make sp. space have a finite no. of pts. in it, i.e. $\sum x = 1/1/m$

Let $N(x, k)$ be th. prob. ~~that~~ x will have k kids.

Let D_{xy} be th. ext. prob. from x to y

- usually $D_{xy} \neq D_{yx}$. [the it may in general, due to Q.M. symmetric matrices]

$$P_{n+1}(x, k) = \sum_{l_1=0}^m \dots \sum_{l_m=0}^m \left(\prod_{i=1}^m P_n(i, l_i) D_{i, x} \right) \quad \text{for } \sum_{i=1}^m l_i = k$$

this is wrong; $N(x, k)$ has been forgotten.

This is a summation over all possible ways in which one can have transitions to x giving a total of k individuals at x , i.e. l_i kids from ~~all~~ i going to x , with prob. $D_{i, x}$, and $\sum_{i=1}^m l_i = k$, so that the total transfer is k .

A "good" metric is one in which D_{xy} is small if $\sum_{k=1}^m |N(x, k) - N(y, k)|$ is large.

Z 13-16

- The important problems, then are
 - For a given metric, and environment and unit conditions, will the species die out or expand? — vary all 3 parameters.
 - How does changing environment with time affect the above? — can simulate this effect by a constant environment and a "perpet" metric?

One can try a few simple distributions for metrics, environments, etc.

Unless I think of some very good ideas, this seems to be about as far as one wants to go at present. The tie-in with the "Black Rap" problem is made in Z 11-20.

Metric $\equiv D_{xy}$
Environment $= N(x, k)$
Unit.conds $\equiv x_0$



The tie-in between "Evolutionary T.M." and the "Black Box".

On the bott. of Z 12 we expressed the various steps in a study of how the "Evolutionary T.M." works.

Now, using Black box methods - i.e. the idea of an a-priori measure (not metric) on Op. space - functionspace. Op. space \cong space is the space consisting of all possible operator spaces. A possible function on op. space would be $N(x, k)$ of Z_{12-9} - th. prob. that x will have k kids. \cong the space of all possible functions, $N(x, k)$, is op. space funct. Space \equiv OSFS.

If x is a discrete variable over m values, then $N(x, k)$ is a $m \times k_{\max}$ dimensional vector.

Any function on this $m \times k_{\max}$ dimensional space is a funct. on OSFS.

Z 12-22 A "good Metric" will, optimally, I think, have only one local max. (or min). Probably, then, one ~~metric~~ should try to find a transformation from the metric and has, to a better one. Such a transformation does, I think, always exist. I.e. even in the N -dimensional continuous case, one can place all pts. of = goodness near one another. - This involves, in general, discontinuous x funs., but for the "Evolutionary T.M." we have a discrete x operator space, and so this is no problem.

The lesson to be empirically learned from nature is - just what is her metric space? - Also, (to a lesser extent important, because we have some idea of an answer) - what is her "environment" - i.e. "Goodness function" on Operator space?

Most important of all: What measure space is implied by Nature's method of solving her black-box problem? For each measure space there exists a method of soln.

Just what measure space corresponds to Nature's method of soln? There may be no answer to such a problem, since th. unknown measure space methods.

may be characterized by certain procedures that make them "different from the method that Nature uses. If so, ~~why~~ possibly Nature should be improved upon - at least slightly, since the measure space method may, inherently, be the optimum method (not necessarily - the nature probably

Perhaps Analogous - or even more than analogous ^{can be improved upon.} is the case of probability. Suppose one has a problem method of assigning probabilities to any contingency on the basis of any evidence. This method consists of a table with contingency-evidence pairs assigned probability values. Now one can, on the basis of ~~the~~^{relatively} few members of the table, construct a ^{carnap} ~~a~~-prior probability of various states of the universe, and from this, derive the rest of the entries in the table.

But these derived values will not, in general coincide with the true values in the table, unless the "probability evaluation method

by which the table was constructed obeys

.20

~~certain laws~~ — and there is no reason to believe it does. It cannot (I think)

be shown, that the carnap-derived prob., is, in any sense, better than the one in the table; because it may turn out that

~~actually~~ all the values in the table were very good, except the ones from which

the carnap probabilities were derived — these latter ^{values} being selected at random.

— Supposing further that all the "good" values were derived from a carnap set of basic probabilities, but not from the ones in the table.

In such a case, perhaps it is possible to modify all of the probability values so that they approach, more probably, a truer state of affairs. Suppose, for instance, that one could assign an a-priori prob. to each carnap-type contingency-evidence-prob. table, and one had some sort of ideas as to what kind of distortion forces might produce this table, one could work back and find inverse prob. — i.e. the prob. that each carnap-type

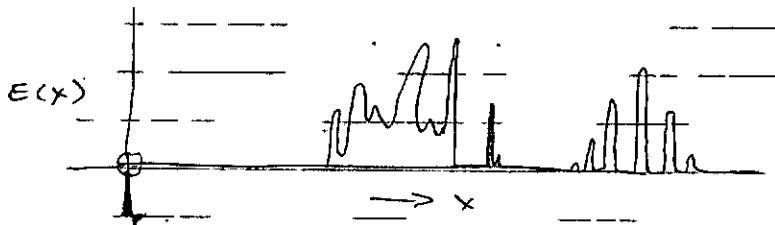
$$F(A \cdot B) = F(F_A + F_B)$$

Table was the original one. This is an interesting problem — Given a prob. eval. method (\equiv contingency evidence = prob. table) to find, from this, the most probable "true" probabilities — if such exist.

216.1

Getting back to the evolutionary T.M.: $N(x, k)$ could, as an approximation, be made a poisson distrib., and it has only 1 parameter (remember variance of Poisson distrib. \propto expected value). $\propto \frac{1}{n!} e^{-n}$ is prob. of occurrence of n ~~occurrences~~^{kids}, if n is the expected no. of kids.

Suppose $E(x)$ where $\bar{x} \leq k N(x, k)$ is the expected no. of kids at x .



Let $g(x)$ be the width of the gaussian distribution of mutations.

Then we have a new $E_{\text{c}}(x)$ which is the convolution of $E(x)$ and $g(x)$. $E_{\text{c}}(x)$ is a more smoothed $E(x)$ and is more useful than $E(x)$.

Suppose each x had exactly $E(x)$ kids each time. Also suppose the x scale is distorted so that the variance of $E(x)$ is constant for each x .

Let $P(x)$ be the initial population density distrib.

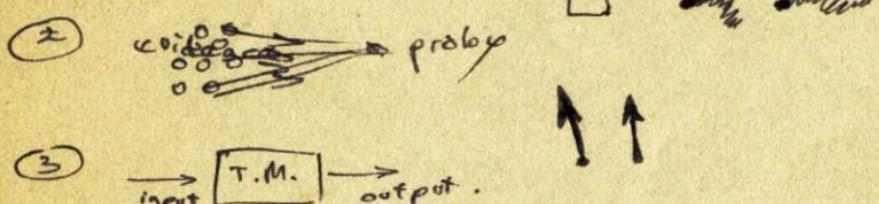
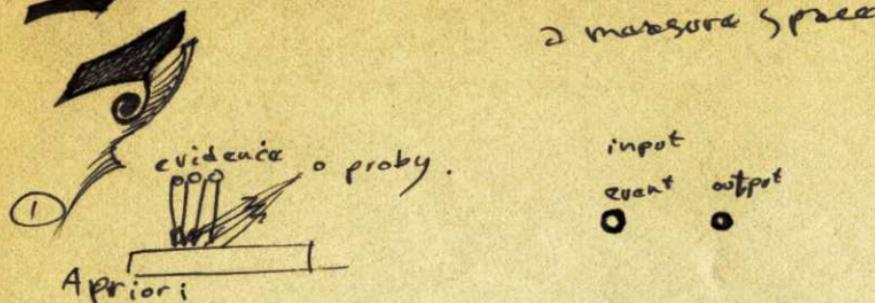
Then proceeding in a naive way, convolution integral:

$$P_{\text{new}}(x) = (E(x) * P(x)) * g(x)$$

$$\approx P_{\text{new}}(x) = \int (E(x) * g(x)) * P(x) dx = \int g(x) P(x) dx$$

Here we have avoided sex completely. This is a very important factor in the Evolutionary T.M. It would be interesting if one could show that the evolution of sex was very probable. Also the presence of sex indicates something about the matrices.

Relatedness. (A metric space further than a measure space.)



optimization of reproduction proby.
can be done if p.e.m. is a rel.
 but existence of p.e.m.
 is an a.l. assumption.

(+) However, a human being is able to solve problems other than the problem of reproduction — so that perhaps the elementalism suggested by p.e.m. is not a bad one.
 i.e. it would seem poss to learn from ~~a~~ human being ^{behavior}, how to solve problems as a goal. we do not have to be so non-el. as to allow only to prod. to be a goal.

~~B~~
 Alteration of moves between T.M. and env.

{ Preliminary concept of "Good"
 which is elaborated upon as time progresses.

{ Concepts of Similarity of inputs.
 Prelim. concept which is elaborated upon as time progresses.

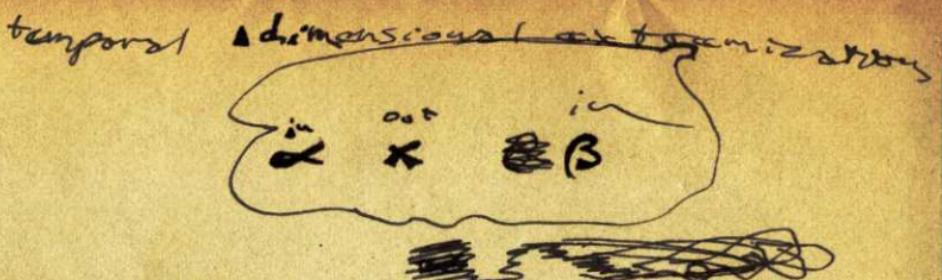
~~Inputs~~
 2 events are ~ when they are followed

~~Inputs~~ & ~ outputs

make them be followed by ~ inputs.

~~Inputs~~ ~ (and conversely about ~ outputs)

At first 2 inputs or 2 outputs are ~ if they are usually followed by ~~good~~ or bad.



$$\int_{-\infty}^{\infty} \exp\{-\frac{1}{2}(S(t) - R(t))^2\} e^{-i\omega t} dt$$

Basic idea of learning is to know what strings of stim-resps. demand the same responses.

Generalized similarity:

if 2 situations containing many stimuli and responses, are similar in detail, except event η_1 = (stim. or resp.) of situation 1 and η_2 of situation 2 } ~~and~~ η_1 and η_2 are not a \sim or dissimilar - they have simply not been evaluated, then η_1 becomes \sim to η_2 .

In addition to \sim and α for every ~~one~~ SS or RF pair one must have a degree of certainty of each value, to tell how much weight it must be given. ~~when mixed~~ with other evidence.

A ^{general} device that is less shock and more general than ≥ 20 (top) would be a mechanism to keep T.M. non-ai. I.E. it would watch to see that the end-goals were being pursued, and that T.M. didn't become "perverted" in over emphasis of ^{unimportant} sub-goals.

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March 1, 1952

Z 15 - 4 This is a very important point, i.e., the best probability method is probably not the "normal" one. ~~Normal pr. & v. meth's are ones to which a measure space can be assigned.~~ In the past, I have assumed that a p. & v. meth. using a metric space (like the one used by Janes) was a "normal" one — but this is now (it appears) extremely unlikely.

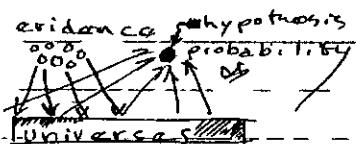
(At the present time, an expedient course of action, would be to investigate some P.E.M.'s (prob. evaluation methods) to find out "how" "abnormal" they are, or just to get an idea of what they look like. Consider P.E.M.'s implied by variations on Wiener's "affectional tone mechanism". It seems, at present, that the ~~"AVERAGE"~~. Looking at P.E.M.'s from the pt. of view, of "Normality" is sterile — that examination of language and general scientific methods and learning theory, in attempts to get a metric space, would be better.

The idea of a metric space. This is a space of events or abstractions, such that if 2 events are close to one another, then they are also "close" in "goodness" value. I think that many living beings use a metric space in which space and time proximity imply closeness of G (goodness = G) value. For metric spaces, it is best to have one with only one maximum point for G — i.e. no local ~~maxima~~ maxima or even minima. (Usually, of course, this is impossible.)

Review:

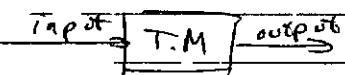
- ① A "normal" prob. is one in which it is possible to assign a priori probabilities to all the possible states of the universe. When evidence is offered, certain of these universes become impossible. Then the probability of any statement is simply the sum of the ^{renormalized} probabilities of all universes in which that statement is true.

One starts with the a-priori universes. The evidence modifies the prob. distribution, and from th. new distrib., one can calculate prob. of



(2) An "abnormal" p.e.m. is one in which the prob. assigned to any evidence, hypothesis pair cannot be derived from any such a-priori universe probability.

(3) The problem of th. human T.M. is to maximize reproduction rate. If a p.e.m. is available



this problem can be

optimally solved. However,

~~(3)~~ that a p.e.m. is needed for ~~area~~ the solution of such a problem is an elementalist assumption, and is probably untrue.

(4) That the human T.M. can solve problems other than maximization of reproduction rates indicates that the elementalism of (3) above may be OK. Also, that one may be able to learn how to solve simple problems by observing human behavior.

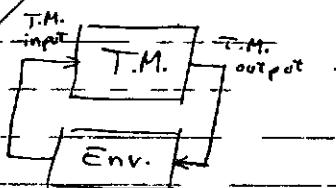
Investigation of th. following kind of T.M.

I think this is similar to Wiener's Affective tone mechanism.

Env. [def.]

(1) Alteration of ^{digital} moves between T.M. and Environment.

(This may or may not be a simplification but it isn't too closely related to the assumptions which follow)



Note : In th. discussion to follow
All inputs and outputs refer to
T.M.

(2) T.M. starts with some rather vague, loose concepts of "good" and "bad". They assign a number to each possible input. They may be vague, and loose for one or both of 2 possible reasons:

2) The they cover all possible inputs, the metric space they define is largely on the basis of proximity in time and/or space. I.E. 2 inputs have rather close goodness values (G values) when they are similar in color, shape, sound, position, texture, etc. "Similarity" here, is largely on the basis of physical science meaning, and ^{bio-physics} discrimination threshold - discrimination thresholds do define a connectivity of a space in a topological manner, and do give an idea as to how to construct a metric.

These values are "loose" because they are usually easily changed.

Some G values like satisfaction of hunger, physical pain, etc. are not loose but tightly fixed.

The initial G function is the best that can be made, due to lack of knowledge about the specific universe that T.M.I.S. to be born into.

To review: The G values are assigned first to such things as satisfaction of basic body needs, pain, etc.

Also G values are assigned to general things like temperature, texture, loudness, etc. Note that the latter qualities are general so that "temperature" means average body temperature. G values would be the same for {warm hand and cold foot} and {cold hand and warm foot}

(3) Elementary, preliminary

② above. At first 2 inputs are similar if they have close G values. There is some differentiation, however, even at first, since certain inputs ~~them~~ may have equal G's, yet may demand different responses (outputs). E.g. Hunger of infant plus presence of mother's ~~test~~ in his mouth may produce sucking response; General body warmth and comfort without hunger, may ~~be~~ have equal G, but will produce no sucking response, therefore must be recognized as different situation.

So at first "similarity" is on 2 bases:

- 2 situations are similar if they have close G's.
- " " " " " demand the same response.

Actually 2 may, in a sense, be regarded as a more important criterion in later life. [I think "similarity" is more important than G. G is an aspect of similarity.]

These are preliminary stages of T.M. T.M. "learns" in later life by modifying its concepts of G and of similarity.

In general, if 2 inputs are similar (demand the same response) then they usually have close G values. But if 2 inputs have similar G's, they need not be similar ($\approx \sim$ similar).

How learning modifies:

- 2 inputs, A and B are $\approx \sim$, if input A is followed by ^{output} α and produces input α

" B " " " " β " " " b

* $a \approx b$ and $\alpha \approx \beta$

Then $A \sim B$

- 2 outputs α and β are \sim if:

A is followed by output α and produces input a

B " " " " β " " " b

$A \approx B$ and $a \approx b$

Then $\alpha \approx \beta$

Notation: Capital Latin letters are inputs.

Greek " " outputs.

$A\alpha \rightarrow A$, means A is followed by α producing A, input.

An OLD PROBLEM of the Affectionate T

Mechanism:

- If $\exists \alpha \rightarrow A \alpha \rightarrow B$, and B is Good, then A becomes Good.

The trouble was, that this led T.M. to seek A as a subgoal in itself. Also we can find instances of this in human beings, as a perversion, e.g. the seeking of money as an end, rather than a means; it would seem that such perversions would occur even more frequently than they do, if say, human beings were governed by this mechanism.

I believe that the following solution for the dilemma was proposed when it first came up, or some time after, or now for the first time. This resolution is largely ad-hoc, and is ~~designed~~ to fit observed human behavior.

If $A \rightarrow B$, and B is good than A becomes good. Now suppose B is very important, permanent good, like satisfaction of hunger. Then A will become good and retain its good only by virtue of $A \leftrightarrow B$, so that if $A \not\rightarrow B$ with reasonable frequency, then the "Good" of A begins to drop off.

If, however, the good of A becomes very firmly established (in infancy of a human being, say), it can become a perverted sub-goal in itself, so that even when A no longer produces B , A retains its goodness.

Also - its rather non-el. way of doing it since ~~and~~ for more general learning, as well. X 14.36

This hypothesis has a little ad-hocness, when one considers that it works for conditioned responses, since ~~and~~ for more general learning, as well. X 14.36

~~B (which is the ultimate goal) controls the A.T., not~~

Remember, however, what Goldstein said about A.R.S. — Yet even

A) They may be a parody of learning, they may preserve many essential features of it.

An important question in any learning theory is not if it explains C.R.S. with their many trials, but does it explain rapid learning due to insight? — i.e. Gestalt — the "Ah Ha phenomenon"

A preliminary skirmish with the Gestalt problem, before we go out to lunch.

(1) As T.M. tries to decide what "move" to make, it evaluates goodness (G) of each pass more on the basis of "similar" moves in the past. When it suddenly finds a move of very great expediency and G value, this may be "Gestalt".

(2) Gestalt may be an entirely separate phenomenon from ordinary learning. This, of course, makes the mechanism very ad-hoc.

The Manner of operation of T.M., then, is as follows.

At time t , T.M. has complete recall. It has ~~MINIMUM~~
 $D(x_1, x_2, t)$ the distance between outputs x_1 and x_2
 at time t . It has $\Delta(\varphi_1, \varphi_2, t)$ the distance
 between inputs φ_1 and φ_2 at time t .
 It has $G(\varphi, t)$ the goodness of input φ at time t .

Now T.M.'s next move, $x(t+1)$ will be made by evaluating
 the expected goodness of all possible moves. This will be
 done by assuming all possible sequences of T.M. - Env. moves, which start
 with x_t and extend, ~~ways~~^{next} to moves into the future. The
 a-priori probability of ev. responding ~~in~~ ^{any} way can be gotten
 by using ^{entire} previous history of T.M., plus the φ and x
 distance functions. Since, in general, each proposed
 sequence has not happened before, one will have to
 weight it probabilistically on the basis of sequences which
 are close to it. The degree of closeness is
 given by the φ and x distance functions.

Now - this determines T.M.'s next moves -
 but how should the φ and x distance functions
 and the $G(\varphi, t)$ function be modified by time?

(perhaps it is unwise to do so, since T.M. already has
 behavior which is based, in a very useful way, upon past experience)

Yet I think ^{time} modification of these 3 functions
 is what gives Gestalt!

The variations of $D(x_k, t)$ and $\Delta(\varphi_1, \varphi_2, t)$
 and $G(\varphi, t)$ with time can be gotten using
 the rules [Z 19-15] and [Z 19-20].

An analysis of this T.M. could proceed in
 the following way. Make the T.M. a black box
 problem, with ~~its~~ env. as the black box, and certain
 ultimate goods and sub-goods in T.M.. Sequences
 of T.M., env. moves, alternations are inputs to the
 black box and outputs of T.M.. Successive outputs
 of the black box are inputs to T.M. and are
 evaluated by T.M.

For problems like material problems, at
 first, the only "good" that T.M. can know is correct
 solutions. Then, as ~~they~~ it works more and more
 simple problems, it will build up a store of sub-goods.
 At first T.M. should be fed simple problems with only a fewness
 moves for answers, so that T.M. will build up a matrix

spaces and "good" spaces.

Along with this mechanism analysis, should proceed a constant reference to psychological data, and intuitive ideas about scientific method.

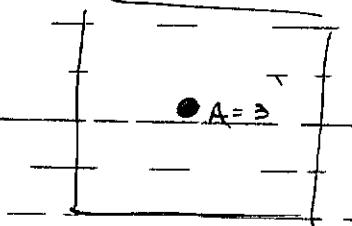
- (05) Note: A less al. idea of goodness and similarity would be for stimulus response pairs or SR's triplets or SRSR quads etc. These would, of course, each have their own spaces, traceable past history and (2) upon simple deductions from assumptions and data in lower order spaces. X^{20.10}

The idea of sub-goals in T.M. is as follows: Suppose one has a black-box problem - i.e. an input and an output representation. In a normal B.B. problem, each output can be immediately given a number, G , and one tries to find an input whose output gives the highest G value. In T.M., however, only a few of the outputs have G values attached to them + and others have G values - but in the form of a probability distribution of large variance.

Again in such a T.M. B.B. problem, one method to maximize G , would be to first fill in lots of the G values in the space of outputs so that one could know in what direction one was going inside the box.

Even with the initial set of G values given, the P.B. may be a hard one to solve, if there are several local extrema.

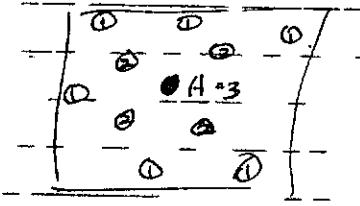
Suppose one has a 2-dim. B.B. (fig. 1)



Say there is only one point, small area, A , that is "good", and one must try to find it by choosing pts. in the box at random.

One would have a rather small fig. 1 prob. of finding it.

Now, suppose that in addition

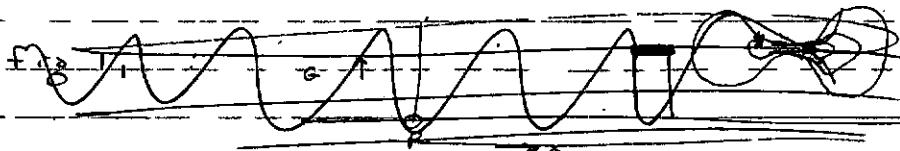


to A , there are other small areas with various G values as shown in fig 2. Then, some random guesses will not give A , but they will

give some idea as to where A is.

Now, say that in addition to the labeled pts. in Fig 2, one has an idea of similarity so that each of the areas shown with a "G" label can be smeared out - but that the parts of the smear outside the original area have variances > 0 in their proby distribs.

Then in 1 dimension:



x is the 1 dim. The $p(x)$ is in the paper, and is the proby density of x having a certain 'G' value. Contour lines of $= G$ are shown.

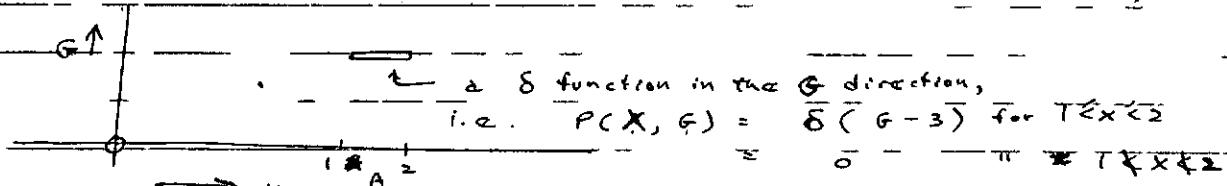
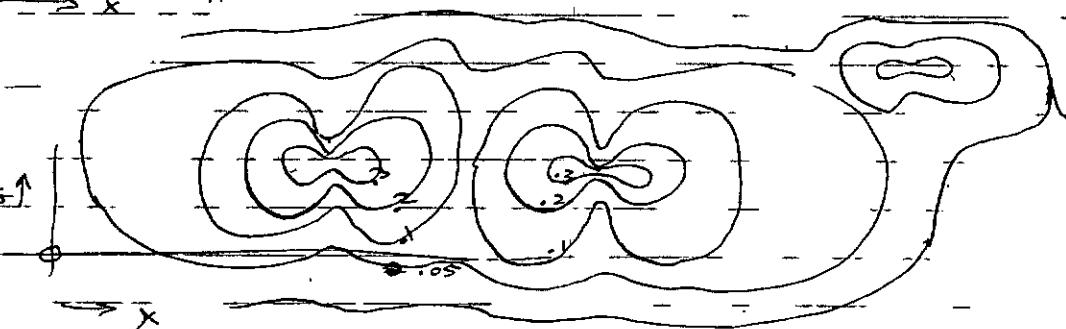


fig 1'

fig 2'



In such a case of smearing, one has even more probability of getting useful info from each trial point.

So one may have as a sub-goal - the realization of a ~~good~~ method to recognize when a pt. is near a really "good" area. The criterion is that these estimates of "good" should be approachable by instruments in leading one toward one's ultimate goal.

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