Applying Machine Learning to Predict ɛ3ɛ4 Gene based on Wayfinding of Sea Hero Quest Game and ACE Cognitive Exam

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The Apo-E gene is located on the 19th pair of human chromosomes. People with an ɛ4 (ɛ3ɛ4) type gene have a 3-5 times increased chance of getting Alzheimer's disease (AD), while the normal human gene is $\varepsilon 3 \varepsilon 3$.

"Sea Hero Quest" (SHQ) is a game that can detect Alzheimer's disease early. According to Dr. Gillian Coughlan's paper(Coughlan, 2019), it was found that the average way-finding distance of ɛ3ɛ3players is lower than ɛ3ɛ4 at the wayfinding game scenes specially in Level 6 and Level 8, and the routes are also relatively concentrated than $\varepsilon 3 \varepsilon 4$.

Addenbrooke's Cognitive Exam (ACE) is an extended cognitive screening technique that including five cognitions: attention, memory, language, language fluency, and visuospatial skills. It had often been used to evaluate AD.

The Rey Complex Figure Test (RCFT, Proposed by Meyers, 1995) consists of a replication test and one or more recall tests . The experiment requires candidates to draw a copy of the graphics presented on the table in front of them. It includes a 3-minute recall trial and a 30-minute recall trial, or an identification test. It was used to detect brain disorders such as Alzheimer's disease, traumatic brain injury, schizophrenia, obsessive-compulsive disorder, Huntington's disease, bipolar disorder, and epilepsy.

This research hopes to combine the SHQ L6, L8 levels way-finding distance, ACE five cognitive items, RCFT test and age to build a neural network model by using SPCN5.0 (Ye, 2009), This work doesn't measure genes directly, but can predict the possibility of getting ɛ3ɛ4 gene which may increase chance of getting AD.

Through Dr. Gillian Coughlan's paper (Coughlan, 2019) 59 volunteers' experimental data. We divide these 59 people into 51 people for training and 8 people for verification. And we use "age sorting" to make 5 different combinations of total 59 data by using stratified sampling method as following table:

mbination	I	II	III	IV	V
mbering	1 · 14 · 30 ·	2 · 11 · 22 ·	6 · 14 · 18 ·	1 × 11 × 30 ×	2 · 14 · 19 ·
To do	34 * 38 * 42 *	25 · 31 · 34 ·	22 · 31 · 39 ·	34 * 38 * 42 *	26 · 33 · 35 ·
rification	44 • 52	49 \ 57	41 • 54	44 [•] 52	41 • 51

The combination table of " Age sorting" and "Stratified Sampling" Then Input of each combination is persons' age, L6, L8 ACE-total and RCFT, Each combination has 8 verification data and the rest 51 is as the training data. We train the model by SPCN5.0 BPN simulation program, then we get 5 verification results.

	Variation	L6+L8+RCFT AGE + ACE	L6+L8+ AGE + ACE	L6+L8+ AGE+RCFT	L6+L8+ ACE+RCFT	L6+L8+ AGE	L6+L8+ RCFT	L6+L8+ ACE
	Combination1	62.5%	62.5%	62.5%	62.5%	75%	62.5%	75%
	Combination2	100%	100%	62.5%	87.5%	75%	75%	100%
	Combination3	62.5%	75%	62.5%	87.5%	62.5%	75%	100%
(Combination4	25%	62.5%	50%	50%	75%	50%	75%
	Combination5	75%	100%	75%	87.5%	50%	100%	87.5%
	Average	65%		62.5%	75%	67.5%	72.5%	

Five parameters is < 75%

Four parameters, Two groups≥ 75% (L6,L8, ACE,AGE) , (L6,L8,ACE,RCFT)

Three parameter, One group $\geq 75\%$ (L6,L8,ACE)

Interestingly, if variation groups $\geq 75\%$, they have L6,L8,ACE factors together!

We go on doing BPN simulation for Prediction., then we get 5 prediction results.

Prediction	L6+L8+RCFT AGE + ACE	L6+L8+ AGE + ACE	L6+L8+ AGE+RCFT	L6+L8+ ACE+RCFT	L6+L8+ AGE	L6+L8+ RCFT	L6+L8 ACE
ombination1	81.3%	81.3%	75%	75%	75%	68.8%	75%
ombination2	81.3%	75%	87.5%	68.8%	81.3%	56.3%	75%
ombination3	68.8%	75%	81.3%	62.5%	75%	37.5%	75%
ombination4	75%	81.3%	75%	75%	75%	68.8%	75%
ombination5	87.5%	75%	81.3%	68.8%	68.8%	62.5%	75%
Average		77.5%		70%		58.8%	

$(L6,L8,ACE,RCFT,AGE) \ge 75\%$

Four parameters, Two groups≥ 75% (L6,L8, ACE,AGE) , (L6,L8,AGE,RCFT) Three-parameter, Two groups≥ 75% (L6,L8,AGE), (L6,L8,ACE)

Similarly, if prediction groups \geq 75%, they have both L6 and L8.

The RCFT seems effect is minor both at variation and prediction!

Now we may look closely on "Line Chart".



There are only two groups that both variation≥75% and the prediction also \geq 75%.

(L6+L8+Age+ACE): Variation Average Value=80.0% Prediction Average Value=77.5%

(L6+L8+ACE): Variation Average Value=87.5% Prediction Average Value=75 %

The L6, L8 and ACE still are the major factors in BPN simulation (but without RCFT) !

As for Age, we can see the red dotted line of three parameters (L6+L8+Age). It's worse at variation (67.5%) but good at prediction (75%). We can hypothesis age is not an important factor in BPN. All we need may be just L6,L8 and ACE-total in BPN.

Conclusion: Combine the Sea Hero Quest L6, L8 levels way-finding distance and ACE five cognitive items, Applying Machine Learning to Predict ɛ3ɛ4 Gene is a potential good way to predict.

PNAS | May 7, 2019 | vol. 116 | no. 19 | 9285-9292 Gillian Coughlan, Antoine Coutrot, Mizanur Khondoker, Anne-

Marie Minihane, Hugo Spiers, and Michael Hornberger(2019)

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