



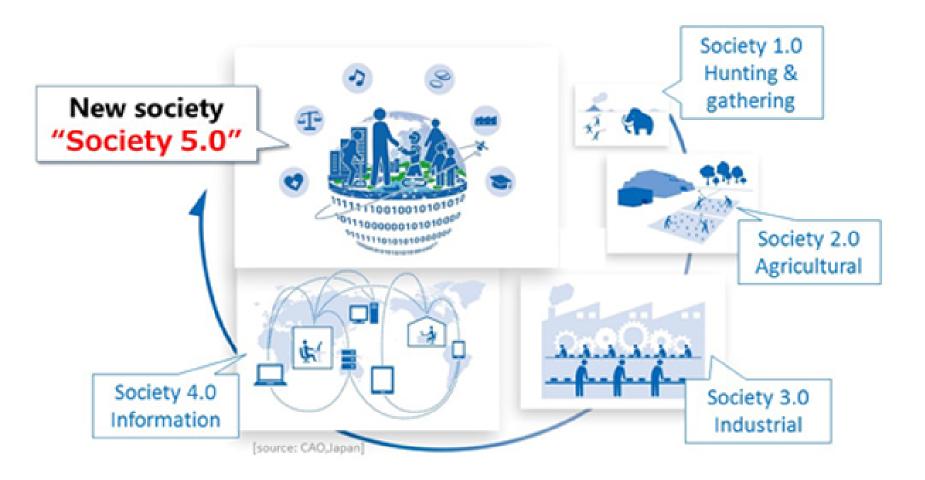


Understanding and predicting disease through AI and mathematical analysis of health and medical data

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SOCIETY 5.0 / PERSONALIZED SOLUTION

The Japanese government is promoting the policy of creating Society 5.0 aims to tackle several challenges by the digitalization across all levels of the Japanese society and the (digital) transformation of society itself.



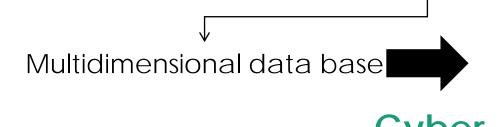
Mission

Precision medicine / Personalized predictive medicine by deep clinical phenotyping



Real life have to be recognized as <u>actual physical characteristics & life course</u>.

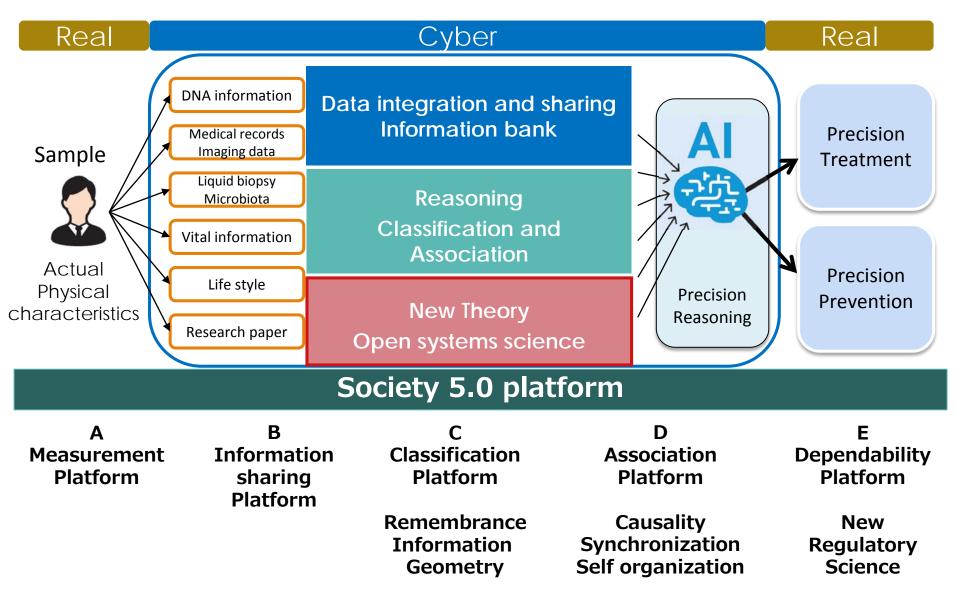
Measurement (Deep Clinical Phenotyping Data)



Mathematically represent and redefine diseases

Goals

■ Platform for deep phenotyping



Machine learning stratification and prediction of ovarian cancer

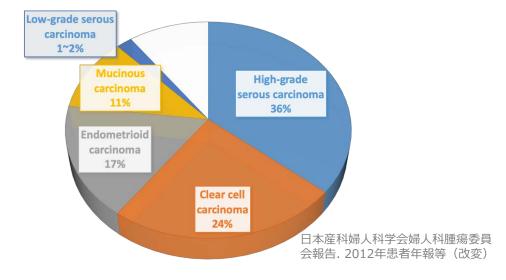
Ovarian cancer in Japan

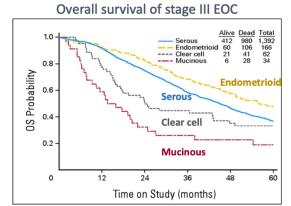
- > 10,048 new patients/year
- 4,745 death/year
- ~50% of patients are found in late stage
- Prognosis is bad specifically in late stage

Stage		5-years relative survival rate (%)
Early	I	87.4
	II	66.4
Late	III	44.2
	IV	28.3

人口動態統計によるがん死亡データ. 2017年. 地域がん登録全国合計によるがん罹患データ. 2014年. 日本産科婦人科学会婦人科腫瘍委員会報告. 2012年患者年報 全国がん(成人病)センター協議会の生存率共同調査(2017年9月)

Histological types of ovarian cancer





Supervised prediction of ovarian cancer

32 preoperative blood markers (CA125, CA19-9, LDH, Fbg, Alb, ···)



Ovarian tumor characteristics (malignancy, stage, histotype)



334 patients with EOC and **101** patients with benign ovarian tumor treated in **the Jikei University** between 2010 and 2017

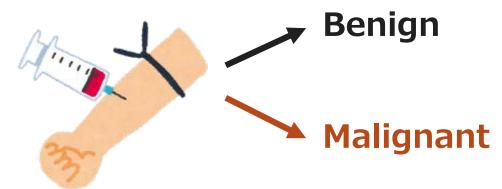
Split datasets into training and validation cohorts (2-fold cross validation)

Train classifiers with training cohort

Check accuracy with validation cohort

Precise preoperative diagnosis based on Ensemble learning

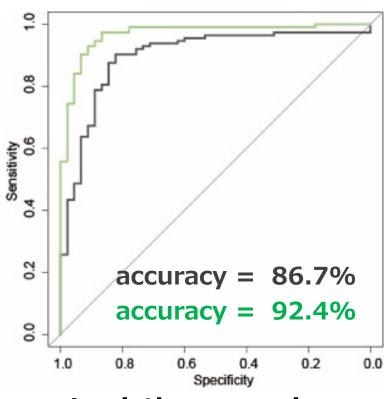
101 Benign tumor334 Malignant cancer



Random Forest

Ensemble of thousands of decision trees

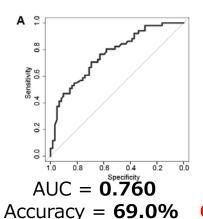
- ✓ Robust to outliers
- ✓ Not depends on type/distribution of variables
- ✓ Consider interaction between variables

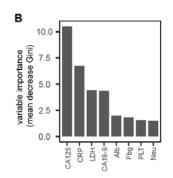


Logistic regression Random Forest

Are known categories really predictable?

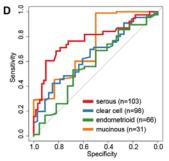
Stratify early stage and late stage

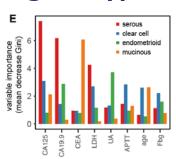




Important factors CA125, CRP, and LDH

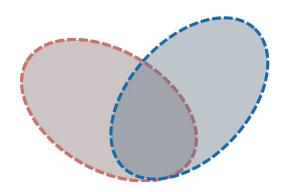
Stratify histological types





AUC = **0.597-0.785** Accuracy = **55.6-96.0%**

Important in **serous CA125, CA19-9,** and **LDH**Important in **mucinous**



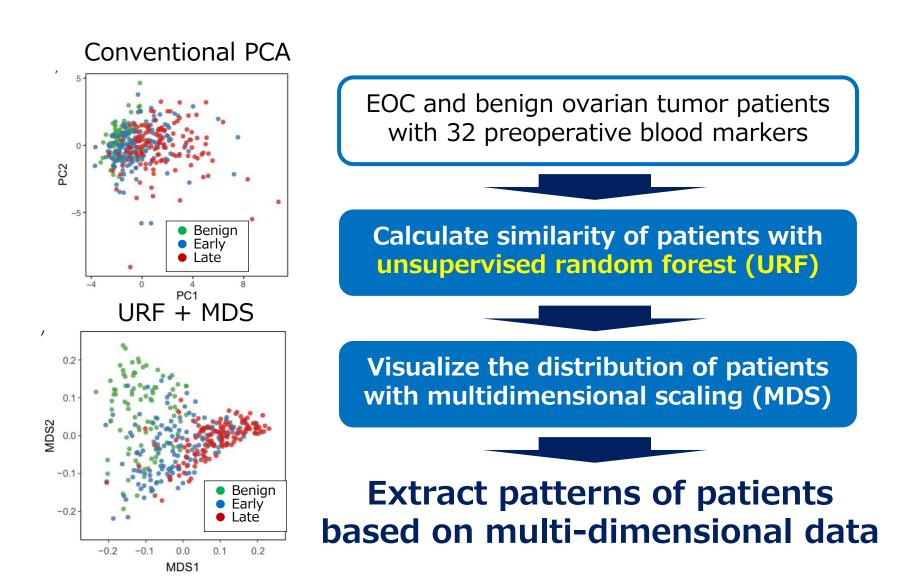
- ✓ There can be overlap between early and late stages
- ✓ Late stage cancer is sometimes mis-diagnosed due to hidden metastasis
- ✓ Supervised prediction cannot find such overlaps



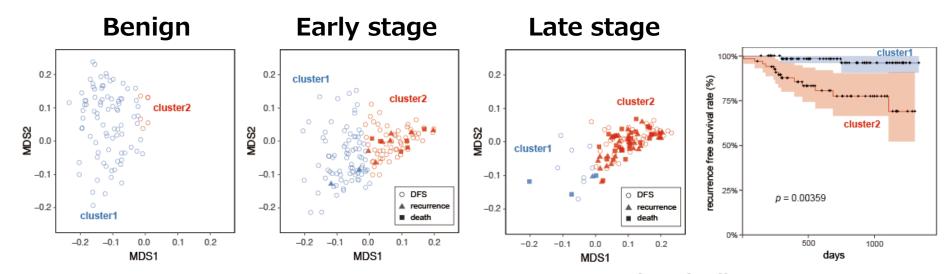
Unsupervised stratification

Kawakami et al. Clin. Cancer Res. 2019

Unsupervised learning to extract difficult-to-recognize patterns



Novel clusters in early stage EOC



Kawakami et al. Clin. Cancer Res. 2019

Early ovarian cancer patients distributed across two clusters

Cluster 1: Benign type

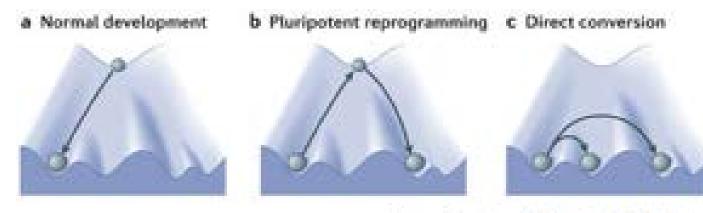
Cluster 2: Late stage type

By using unsupervised machine learning, we revealed the disease clusters that have not been recognized even by clinicians

Landscape modeling of life system



Dr. Tetsuo Ishikawa



Nature Reviews | Molecular Cell Biology

Ladewig 2013
Nature Reviews Molecular Cell Biology

Epigenetic landscape

A concept proposed by Waddington in 1940. The model describes cell differentiation as an analogy of a dynamical system where "state transition occurs as the ball rolls on the terrain".

Data-driven reconstruction of landscape

High dimensional fMRI data

Mean DMN activity

Mean FPN activity

Mean Aud activity

Mean Aud activity

binarize

2ⁿ possible states

110010101011100011

001010001110001110

101101100011000111

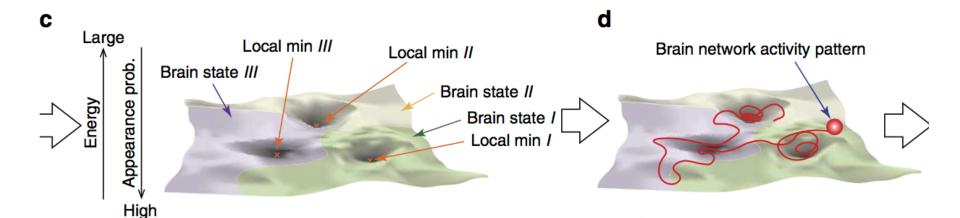
Energy of each state is calculated from observed frequency with Ising model

probability:
$$P(\boldsymbol{\sigma}|\boldsymbol{h}, \mathbf{J}) = \frac{\exp\left[-E(\boldsymbol{\sigma}|\boldsymbol{h}, \mathbf{J})\right]}{\sum_{\boldsymbol{\sigma}'} \exp\left[-E(\boldsymbol{\sigma}'|\boldsymbol{h}, \mathbf{J})\right]}$$

energy: $E(\boldsymbol{\sigma}|\boldsymbol{h}, \mathbf{J}) = -\sum_{i=1}^{N} h_i \sigma_i - \frac{1}{2} \sum_{i=1}^{N} \sum_{\substack{j=1 \ j \neq i}}^{N} J_{ij} \sigma_i \sigma_j$

Characterize brain dynamics

by random-walk simulation

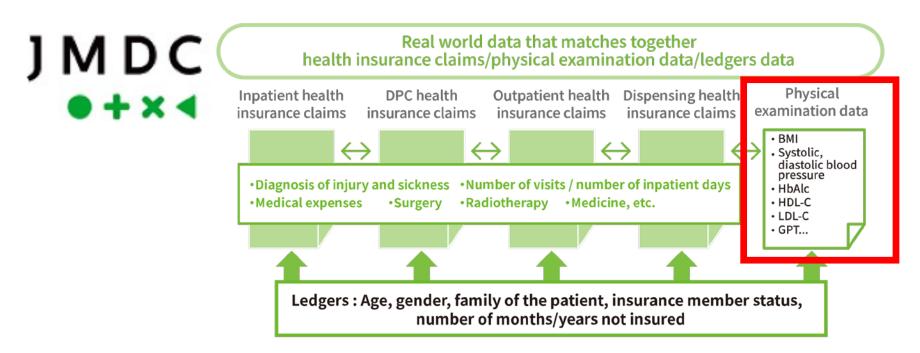


Adapted from Watanabe & Rees (2017) Nature Communications

in the energy landscape

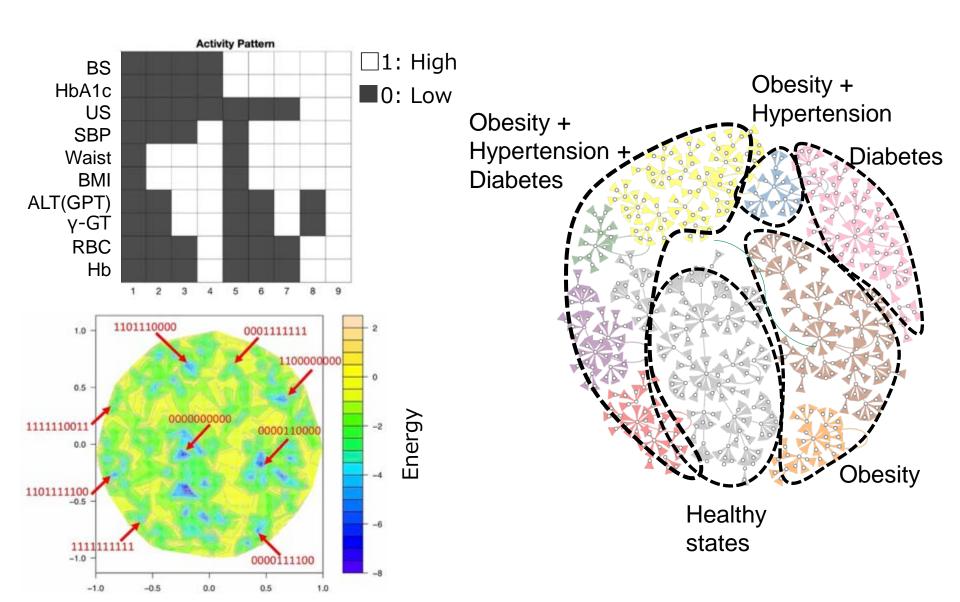
Identify dominant brain states

~2.5M physical examination records

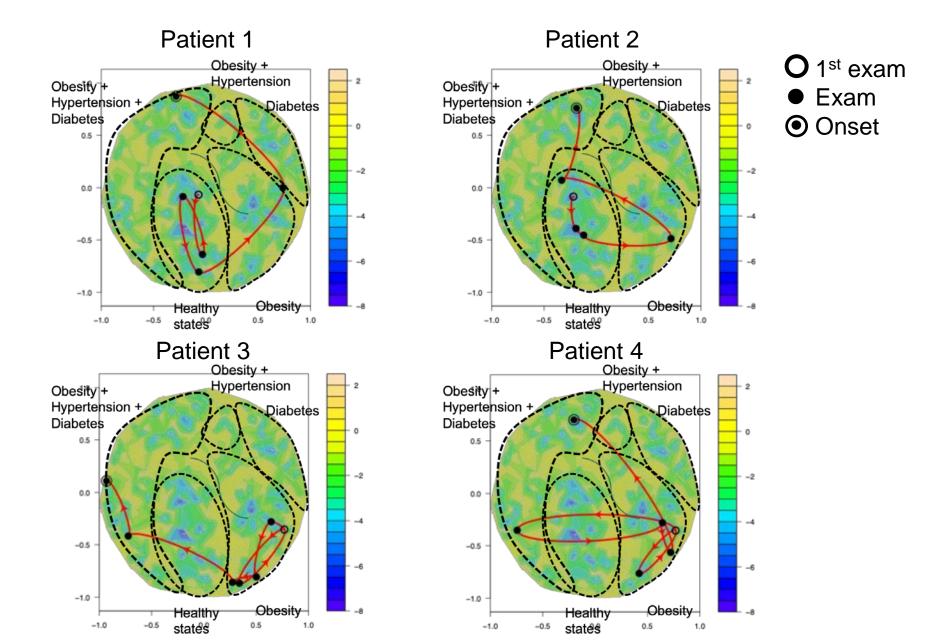


9 健診.c₽hysical examination data 2,581,416 records 683,131 KB			
8 材料.csv	957.608 records	206.889 KB	
7 診療行為.csv	179,979,079 records	35,279,923 KB	
6 医薬品.csv	60,148,880 records	28,782,093 KB	
5 傷病.csv	61,444,578 records	14,967,408 KB	
4 レセプト.csv	26,088,779 records	2,103,235 KB	
3 施設.csv	95,745 records	6,121 KB	
2 患者.csv	331,663 records	18,787 KB	
1 母数.csv	7,163,773 records	405,773 KB	

Landscape of diabetes onset process



Typical diabetes onset trajectories



Key messages

- ✓ Existing clinical classification of disease is not complete. We should re-define healthy/disease states based on clinical phenotype data
- ✓ By using unsupervised machine learning, we can extract difficult-to-recognize patterns that have not been recognized even by clinicians
- ✓ Tracking the onset and progress of diseases may lead to appropriate individualized intervention and treatment

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