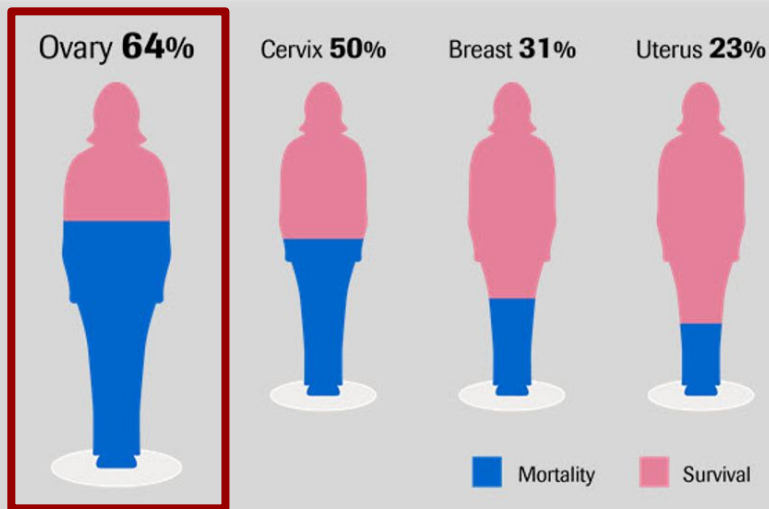


# Platin-drug resistance prediction

Y.-B.Hong

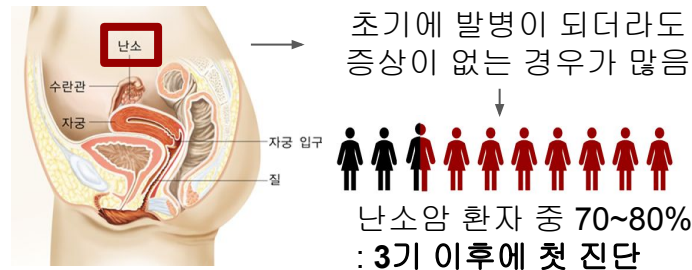
# Ovarian Cancer

## 여성암 중 1위 난소암



(Roche, 2016)

### 1. 난소의 위치적 특성에 따른 늦은 발견



### 2. 높은 재발률

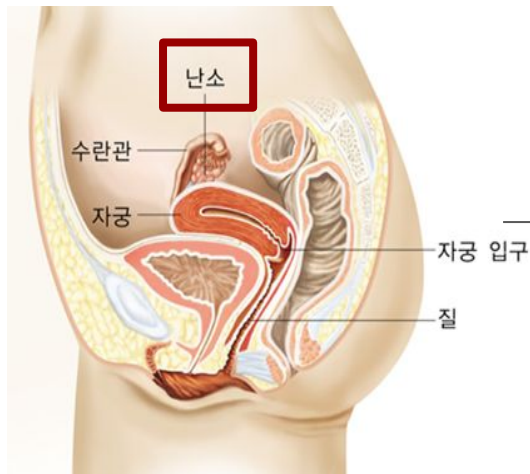
난소암 재발율은 **50-70%**

2년 이내 재발하는 경우가 많다

(대한부인종양학회, 2016)

# Danger of Ovarian Cancer

## 1. 난소의 위치적 특성에 따른 늦은 발견



→ 초기에 발병이 되더라도  
증상이 없는 경우가  
많다 →



난소암 환자 중 70~80%  
: 3기 이후에 첫 진단

## 2. 높은 재발률

난소암 재발율은 **50-70%**, 2년 이내 재발하는 경우가 많다. (대한부인종양학회, 2016)

# Danger of Ovarian Cancer

위치적 특성

복강

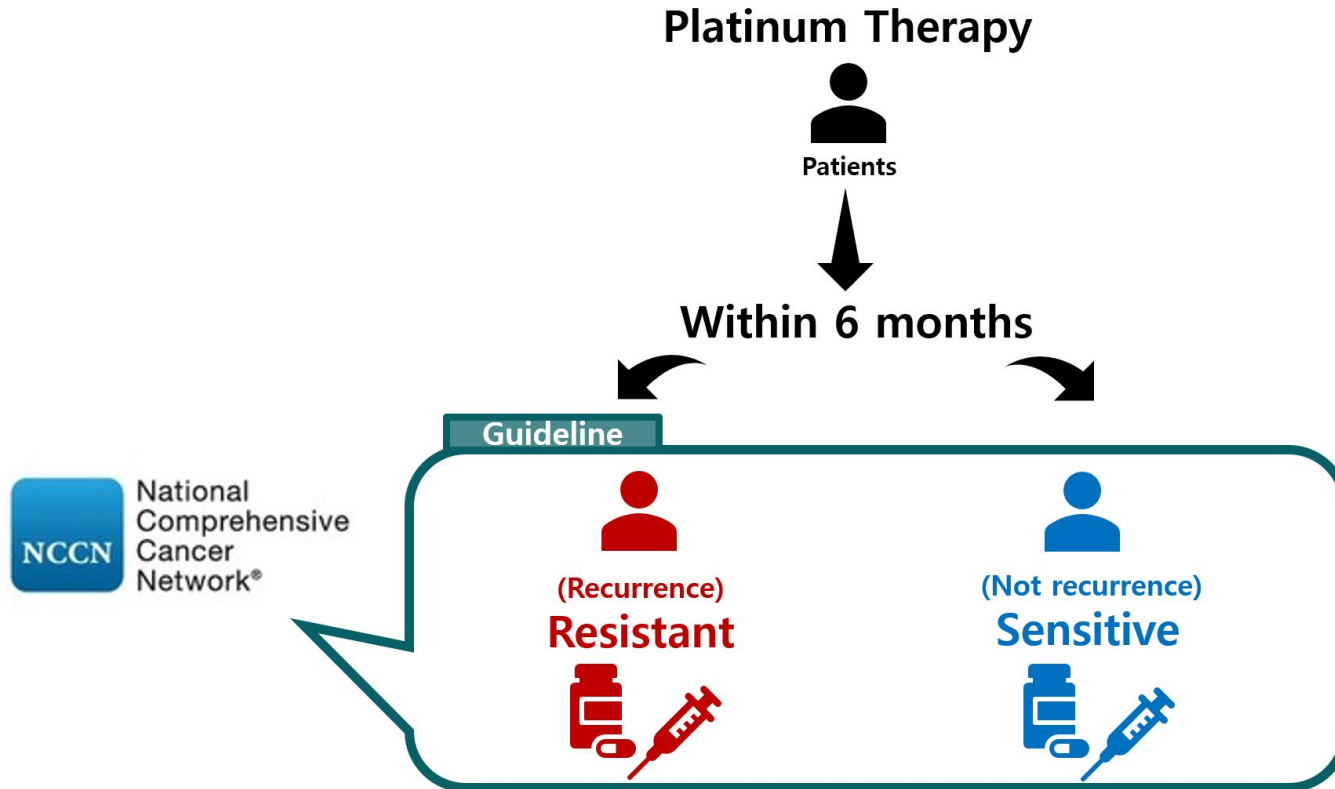
증상을 느끼기  
어렵다

높은 재발률

50~70%

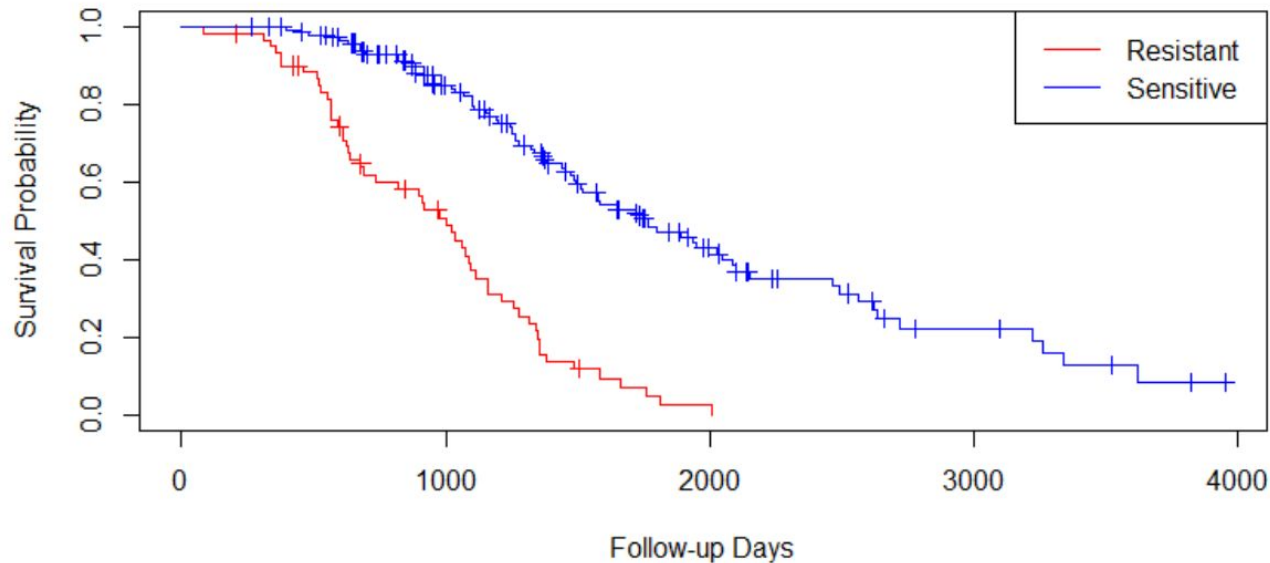
발견되고 나서  
약물치료를 거둬할  
시간이 없다.

# Chemotherapy for Ovarian Cancer



# Chemotherapy for Ovarian Cancer

**Kaplan-Meier Plot for Platinum-resistant and -sensitive Patients**

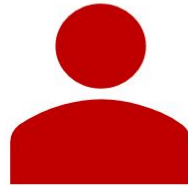


Resistant	Sensitive	Total
97	232	329

# The Goal of our model

Our  
Model

**YOU PLATINUM RESISTANCE**

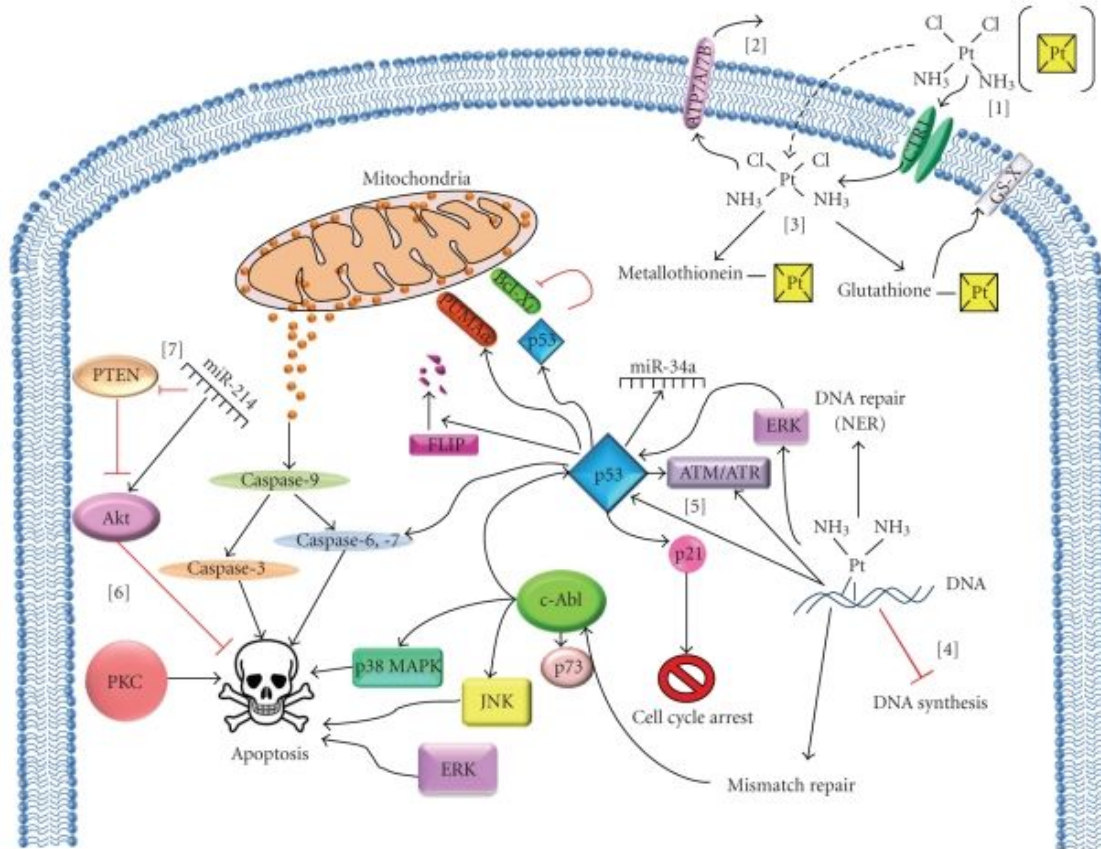


(Recurrence)  
**Resistant**

~~Platinum Therapy~~

**Other Therapy**

# Cause of Platinum Resistance



Pharmacokinetics/metabolic pathway of platinum compound (Basu, 2010).

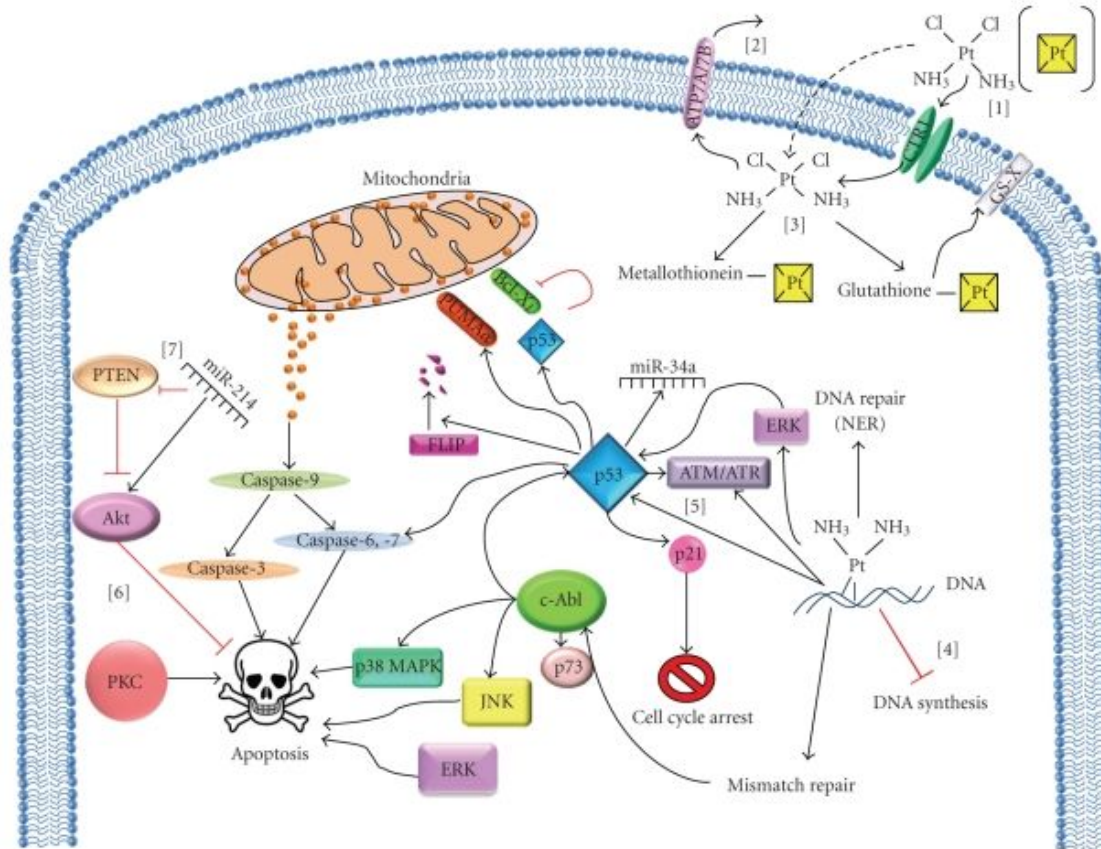
Target therapy for cancer cells:

- DNA damage
- Cell cycle halt
- ER stress
- Decrease in ATPase activity

→ Apoptosis



# Cause of Platinum Resistance



Pharmacokinetics/metabolic pathway of platinum compound (Basu, 2010).

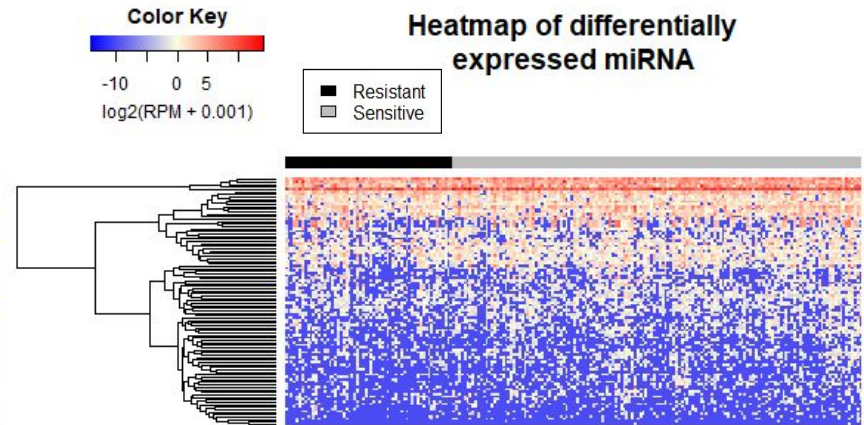
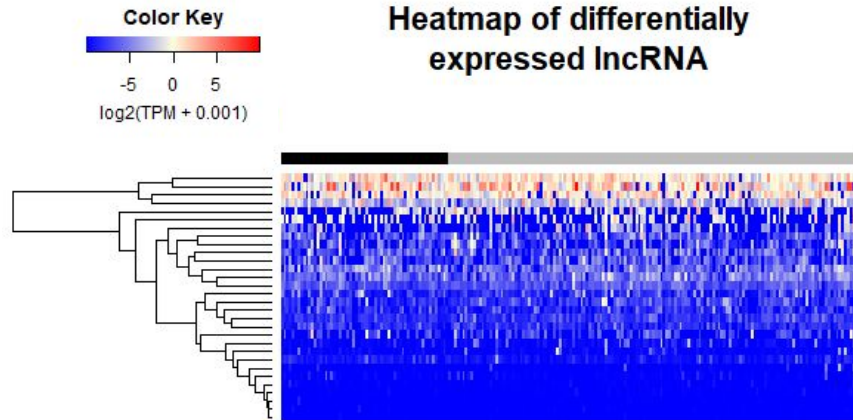
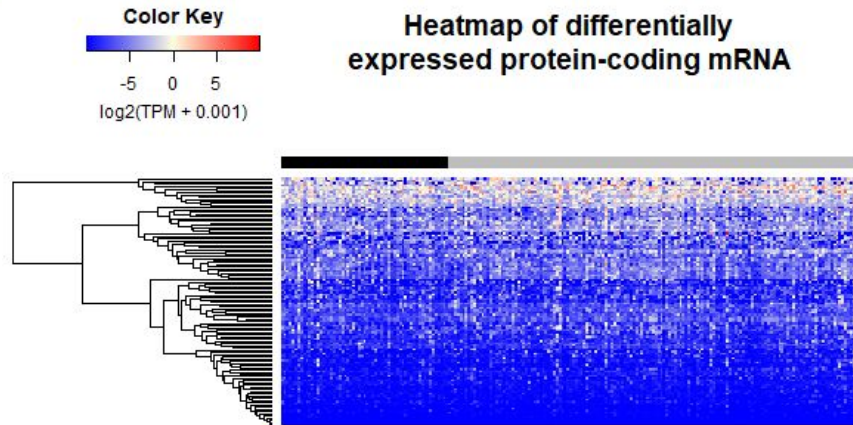
Proteins related to platinum resistance:

- Transporter
- Efflux of anticancer drugs
- DNA repair pathway
- Presence of miRNA and lncRNA

...

Limitation in predicting *in vitro*.

# Cause of Platinum Resistance

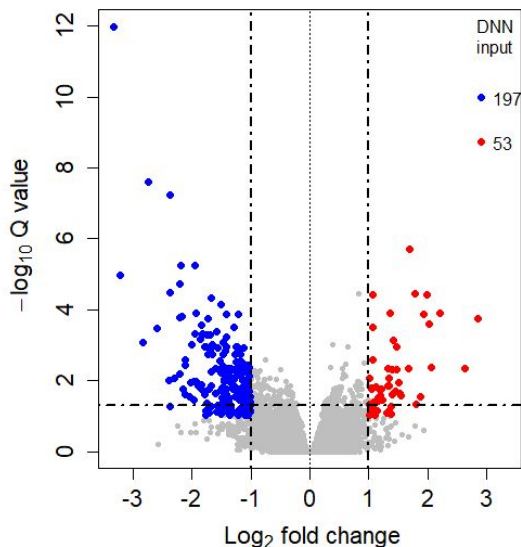


"Differentially expressed":  
A gene which has  $\text{lgFC} > |1|$  and  $\text{FDR} \leq 0.05$  in a differentially expression analysis.

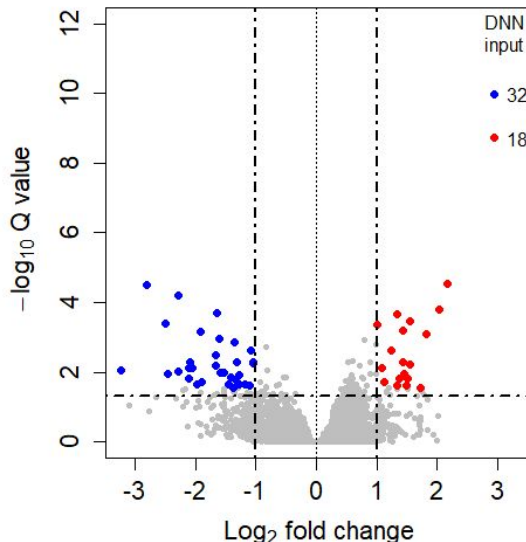
Resistant	Sensitive	Total
55	42	210

# Input for Deep Neural Network

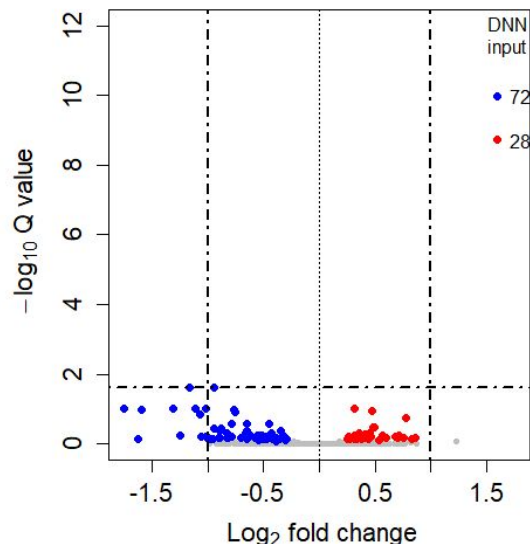
Volcano Plot of Differentially Expressed Protein-coding mRNA



Volcano Plot of Differentially Expressed lncRNA



Volcano Plot of Differentially Expressed miRNA



Patient clinical data:

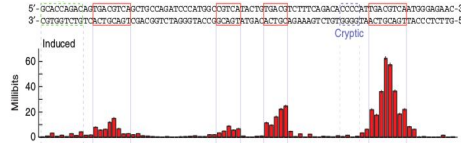
Resistant	Sensitive	Total
55	42	210

Number of input for DNN:

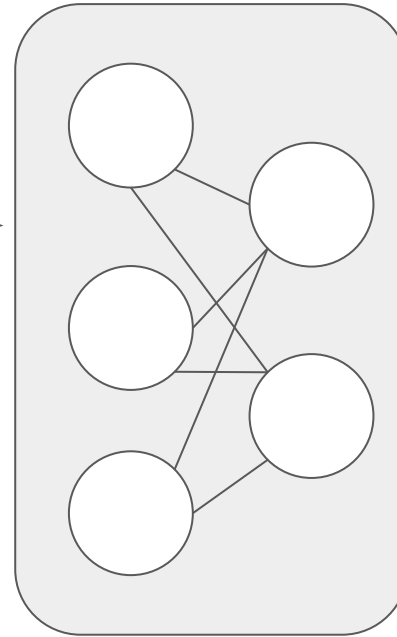
dePC	deLNC	deMI
250	50	100

# Predicting Platinum Status

Backpropagation-based  
Gene expression data



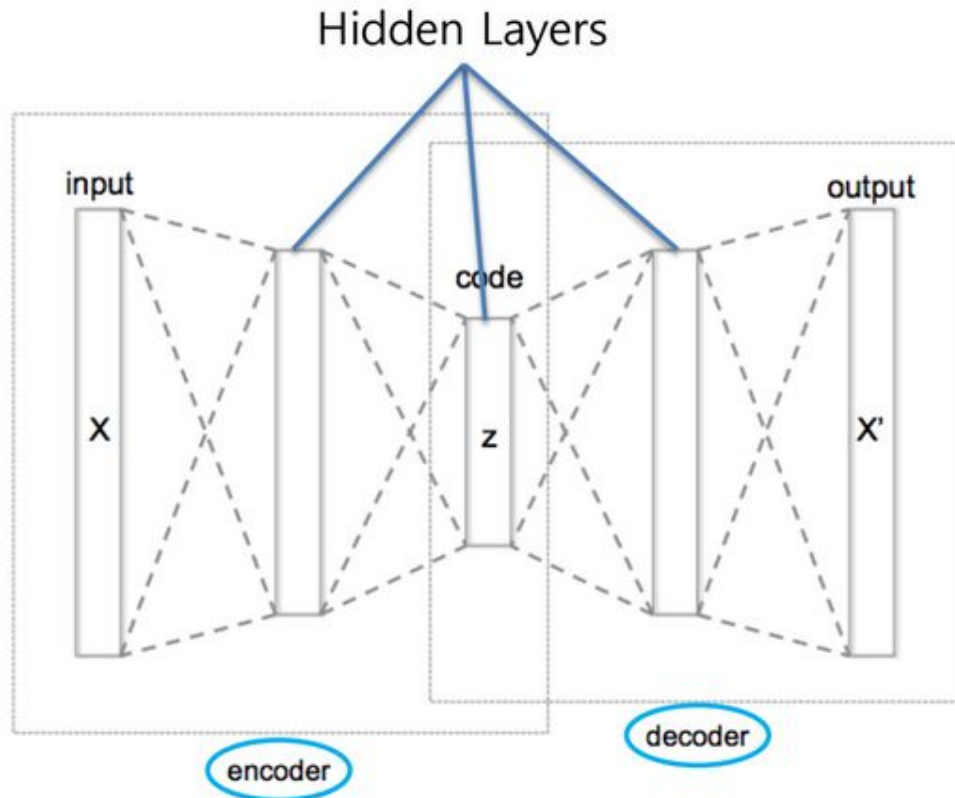
Platinum Status (0, 1)



Predict

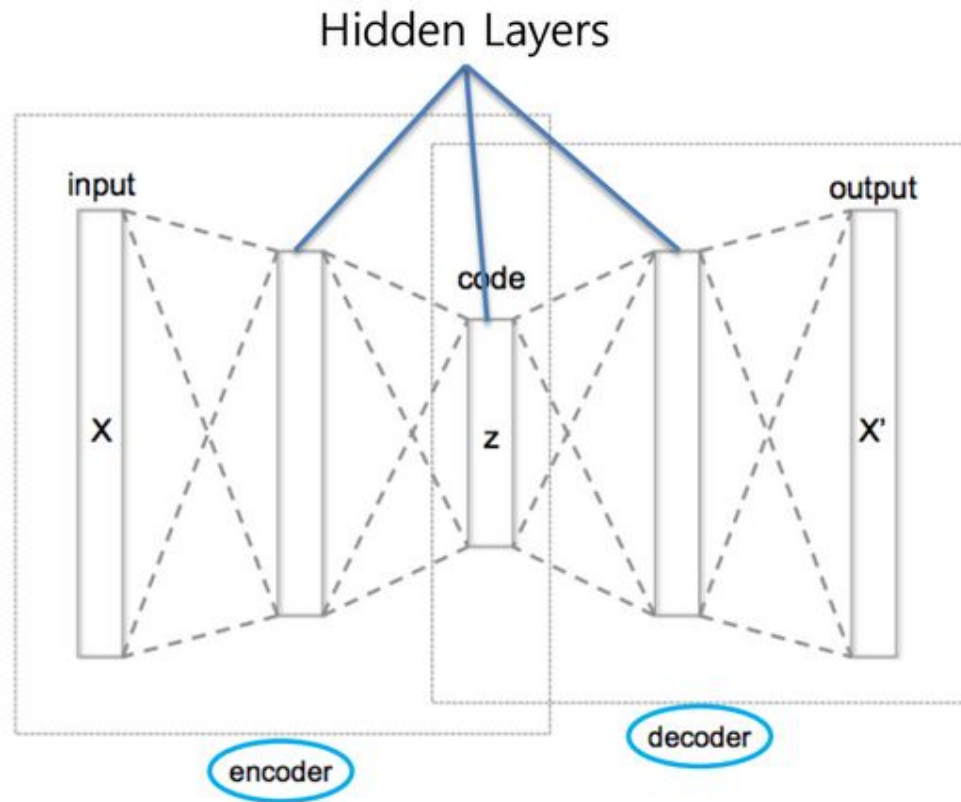
Backpropagate

# Design Model

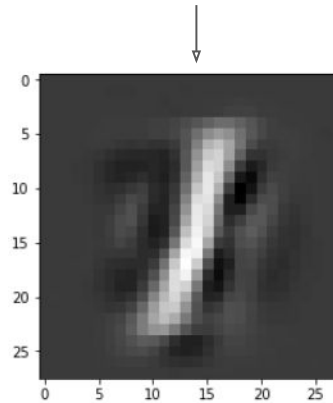


```
class Autoencoder(nn.Module):  
    def __init__(self):  
        super(Autoencoder, self).__init__()  
        self.encoder = nn.Linear(28*28, 50)  
        self.decoder = nn.Linear(50, 28*28)  
  
    def forward(self, x):  
        x = x.view(batch_size, -1)  
        encoded = self.encoder(x)  
        out = self.decoder(encoded).view(batch_size, 1, 28, 28)  
  
        return out  
  
model = Autoencoder().cuda()
```

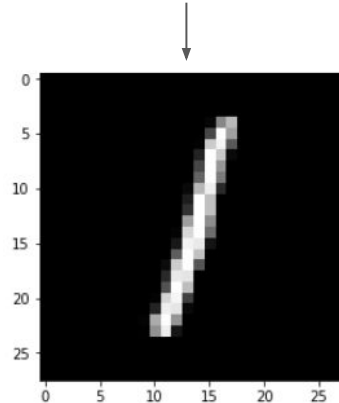
# Design Model - DNN based on Autoencoder



Output Data



Actual Data



# Ensemble - Better voting

High resistance -> 1



High sensitive -> 0



predict  
resistance  
from 1



predict  
sensitive  
from 0



ID	Predict_R	Predict_S
TCGA-29-1769	0.8	0.2
TCGA-61-2097	0.9	0.05
TCGA-23-2078	0.6	0.07

Supported High resistance





# Ensemble - Model performance

Accuracy : 0.8809523809523809

Sensitivity : 0.9230769230769231

Specificity : 0.8620689655172413

AUC: 0.8925729442970822

[[12 1]

[ 4 25]]

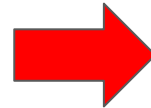
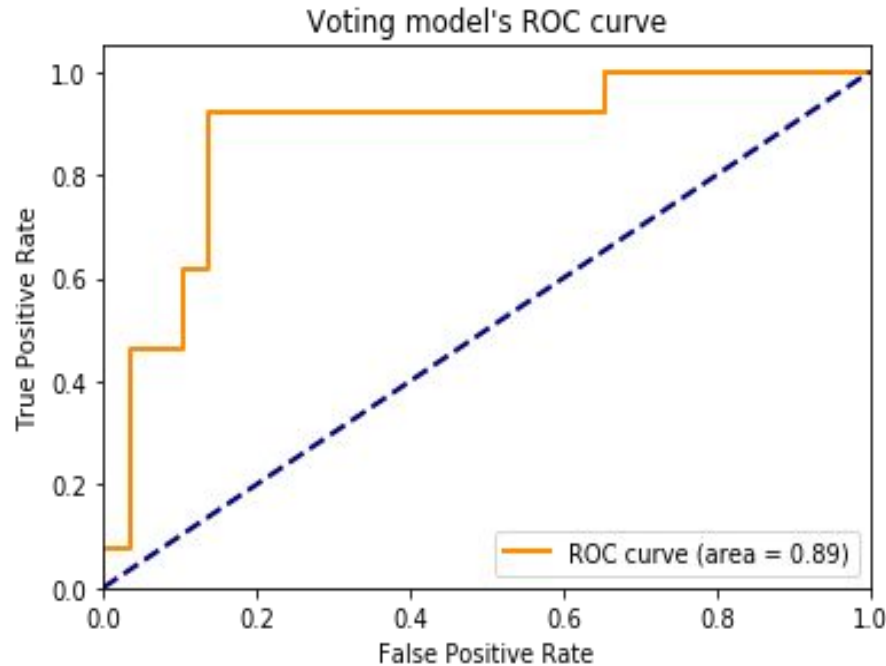
Actual Cases

	Resistant	Sensitive
Resistant	12	1
Sensitive	4	25

predicted Cases



# Ensemble - Model performance and more



Next model?

# Competitor

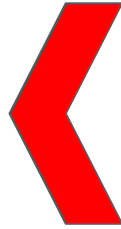
Test(Competitor based on SVM):

Accuracy: 80.0%

sensitivity: 86.7%

specificity: 68.6%

(AUC = 0.776)



Test(Our Model based on DNN):

Accuracy : 88.09523809523809 %

Sensitivity : 92.30769230769231 %

Specificity : 86.20689655172413 %

(AUC = 0.893)

Thank You