# Metric Learning

на примере задачи Face Recognition

# Задача Face Recognition

Верификация (1:1)









Идентификация (1:N)



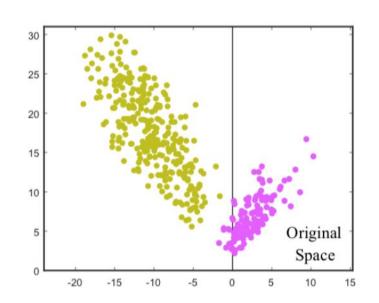


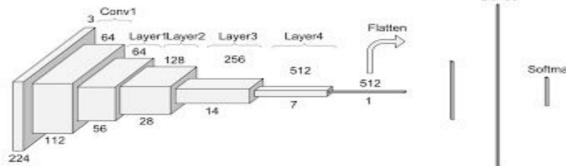


## Классификация:

Стандартная кросс-энтропия:

$$L_1 = -\frac{1}{N} \sum_{i=1}^{N} \log \frac{e^{W_{y_i}^T x_i + b_{y_i}}}{\sum_{j=1}^{n} e^{W_j^T x_i + b_j}},$$







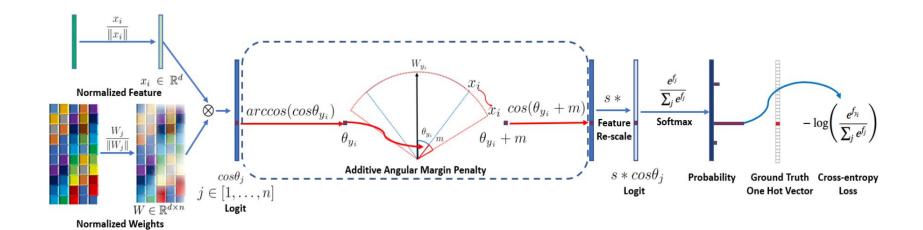
### От Softmax к ArcFace

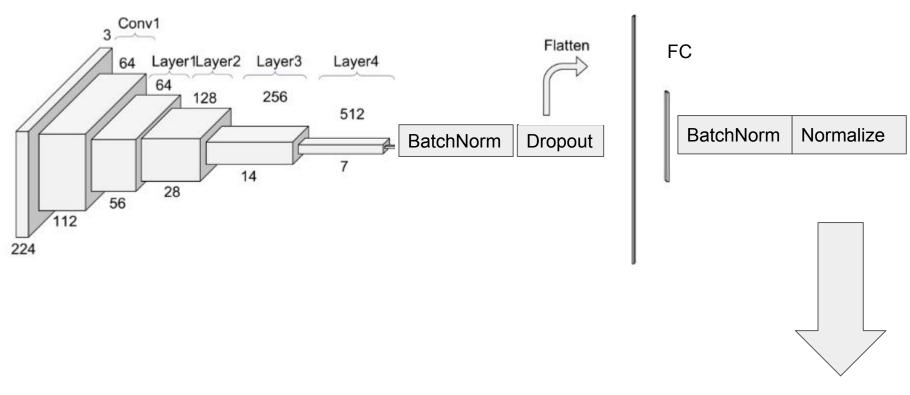
$$b_i = 0$$

$$W_j^T x_i = \|W_j\| \|x_i\| \cos \theta_j$$

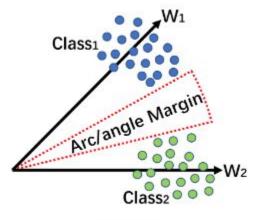
$$L_4 = -\frac{1}{N} \sum_{i=1}^{N} \log \frac{e^{s(\cos(m_1 \theta_{y_i} + m_2) - m_3)}}{e^{s(\cos(m_1 \theta_{y_i} + m_2) - m_3)} + \sum_{j=1, j \neq y_i}^{n} e^{s \cos \theta_j}}$$

### **ArcFace**





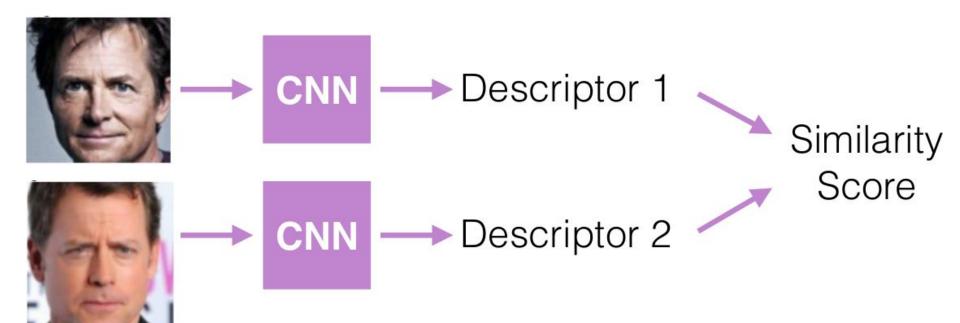
ArcFace loss



(a) ArcFace



(b) Geodesic Correspondence





Overview

### Description

Evaluation

Timeline

Prizes

After centuries of intense whaling, recovering whale populations still have a hard time adapting to warming oceans and struggle to compete every day with the industrial fishing industry for food.

To aid whale conservation efforts,

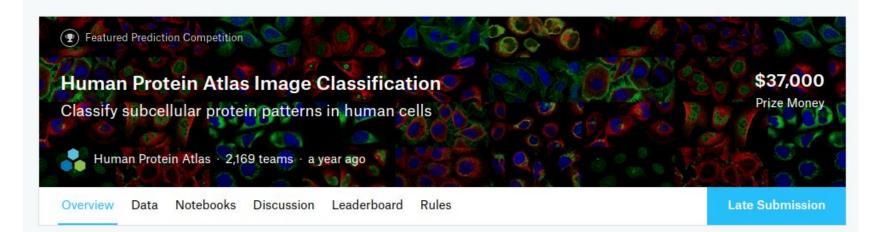
and struggle to compete every day with the industrial fishing industry for food.

To aid whale conservation efforts, scientists use photo surveillance systems to monitor ocean activity. They use the shape of whales' tails and unique markings found in footage to identify what species of whale they're analyzing and meticulously log whale pod dynamics and movements. For the past 40 years, most of this work has been done manually by individual scientists, leaving a huge trove of data untapped and underutilized.

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In this competition, you're challenged to build an algorithm to identify individual whales in images. You'll analyze





Overview

#### Description

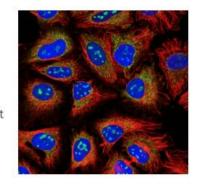
**Evaluation** 

Prizes

Timeline

Special Prize Instructions In this competition, Kagglers will develop models capable of classifying mixed patterns of proteins in microscope images. The Human Protein Atlas will use these models to build a tool integrated with their smart-microscopy system to identify a protein's location(s) from a high-throughput image.

Proteins are "the doers" in the human cell, executing many functions that together enable life. Historically, classification of proteins has been limited to single patterns in one or a few cell types, but in order to fully understand the complexity of the human cell, models must classify mixed patterns across a range of different human cells.



Images visualizing proteins in calls are commonly used for hismodical research and these calls could



Overview

#### Description

**Evaluation** 

Timeline

**CVPR 2019** 

Prizes

Image retrieval is a fundamental problem in computer vision: given a query image, can you find similar images in a large database? This is especially important for query images containing landmarks, which accounts for a large portion of what people like to photograph.

In this competition, Kagglers are given query images and, for each query, are expected to retrieve all database images containing the same landmarks (if any). The competition will proceed in two phases: The 1st phase will use the same test and index sets as last year, while for phase 2 we will release a completely new dataset that contains 700K images with more than 100K unique landmarks. We hope that this release will accelerate progress in this important research problem.

This challenge is organized in conjunction with the Landmark Recognition Challenge (https://www.kaggle.com/c/landmark-recognition-2019). In particular, note that the test set for both challenges is the same, to encourage participants to compete in both. We also encourage participants to use the training data from the recognition challenge (either from this year's or last year's dataset) to develop models which could be useful for the retrieval challenge.

```
class ArcMarginProduct(nn.Module):
    def init (self, in features, out features, s=30.0, m=0.50, easy margin=False):
       super(ArcMarginProduct, self). init ()
       self.in features = in features
       self.out features = out features
       self.weight = Parameter(torch.FloatTensor(out features, in features))
       nn.init.xavier uniform (self.weight)
       self.easy margin = easy margin
       self.cos m = math.cos(m)
       self.sin m = math.sin(m)
       self.th = math.cos(math.pi - m)
       self.mm = math.sin(math.pi - m) * m
    def forward(self, input, label):
       # ----- cos(theta) & phi(theta) ----
       cosine = F.linear(F.normalize(input), F.normalize(self.weight))
       sine = torch.sqrt((1.0 - torch.pow(cosine, 2)).clamp(0, 1))
       phi = cosine * self.cos m - sine * self.sin m
       if self.easy margin:
           phi = torch.where(cosine > 0, phi, cosine)
        else:
           phi = torch.where(cosine > self.th, phi, cosine - self.mm)
       one hot = torch.zeros(cosine.size(), device='cuda')
       one hot.scatter (1, label.view(-1, 1).long(), 1)
       output = (one hot * phi) + ((1.0 - one hot) * cosine) # you can use torch when
       output *= self.s
        return output
```

https://github.com/ZhaoJ9014/face.evoLVe.PyTorch