Machine Learning

Zusammenfassung / Joel von Rotz / Quelldateien

Begriffe

Artificial Intelligence Software, die intelligentes Verhalten imitiert

- ← Machine Learning System, das von Erfahrung/Daten lernt
 - ← Deep Learning basiert auf geschichteten neuralen Netzwerken

Supervised Learning Lernt ann labeled Daten \rightarrow predicts unseen data, Dimensionalitu Reduction

Unsupervised Learning unlabeled Daten → Daten-Struktur lernen und Kunden in Zielaruppen teilen (Recommender Sustems)

Semi-Supervised Learning Mix Sup.+Unsup. → wenig *Labeled* & viele *Unlabeled* **Reinforcement Learning** Belohnungsfunktion → sucht ideales Verhalten für maximale Belohnuna

Regression Problem Label (vorherzusagendes Attribut) ist kontinuierlich **Classification Problem** Label ist kategorial (binary -> Bsp. Ja, nein) **Huperparameters** vom Programmierer ausgewählte Parameter (Bsp. K bei K-NN)

Data Quality Assessment -

- Identify data sources and trustworthiness Check data ranges (e.g. negative salary)
- Interpret statistical key figures Validate plausibility of variable correlations Measure data redundancy • Check for anomalies in sytax and semantics • Explore Null values and duplicate data records

Data Cleaning

Data quality can be improved (although document changes, inform,...):

• identify & remove duplicate data records • replace null values (e.g. median or mean) or delete them (requires large data) • investigate the origins of quality issues

Feat. Eng. - Vector Space Model

Data set consists **only** of numerical data (e.g. color \rightarrow black, red, ...). Categorical data is transformed into numerical data

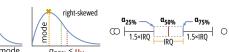
mean μ easy calculate; median robust against outliers; $Var_{smpl.} \frac{1}{n-1}$; $Var_{pop.} \frac{1}{n}$

$$\frac{\text{Var}}{\sigma^2} = \frac{1}{n-1} \sum_{i=1}^{n} (x_i - \mu_X)^2$$
; $\text{Cov}(X, Y) = \frac{1}{n-1} \sum_{i=1}^{n} (x_i - \mu_X)(y_i - \mu_Y)$









Pearson Correlation $\rho(X,Y) = \frac{Cov(X,Y)}{\sigma_X \sigma_Y}$ normalizes ranges: -1=anti-corr., +1=corr

Similariv Measures

Euclidean Distance $\sqrt{\sum_{i=1}^{n} (x_i - y_i)}$ (distance)

 $\label{eq:cosine_posterior} \text{Cosine Distance 1} - \text{CosineSimil.} = 1 - \frac{\sum_{i=1}^{n} x_i \cdot y_i}{\sqrt{\sum_{i=1}^{n} x_i^2} \sqrt{\sum_{i=1}^{n} y_i^2}} \; (\text{distance } + \; \text{angle})$

Manhattan Distance $\sum_{i} |x_i - y_i|$ (along a grid)

Levenshtein/Edit Distance for strings: +1 on delete, add; +2 on change char

Jacard Similarity: $\frac{|X \cap {\mathfrak{a}} \cap {\mathfrak{d}} Y|}{|X| |{\mathfrak{o}} \cap Y|} \to [0,1]$; 1=same

k-Nearest-Neighbor Classification =

Data takes value of majority of k (hyperparameter) neighbors + \circ \circ (optional) weight based on 1/d (distance d). — very slow with lots of data and most comput. is done in classification stage + least data hungry algo, good baseline for other

Normalization

Min-Max: $\frac{x - \min x}{\max x - \min x} \rightarrow [0, 1]$ + interpretation in %, no negatives — no supervised

Z-Score: $\frac{x-\mu_X}{\sigma_X} \to \frac{\mu_{X'}=0}{\sigma_{X'}=1}$ + for (un)supervised — no direct interpr., negatives

Regression Hypothesis & Evaluation Metrics -

Goal is to fit a line or curve-function to a number of points. y_i ground truth, f_i prediction; MAE (Mean Absolute Error), MAPE (Mean Absolute Percentage Error) human readable, MSE (Mean Squared Error) optimized for processing

Model Evaluation Workflow

Workflow: (1) Mix data (2) Split into training & test data (3) train (4) evaluate! NEVER reuse test set!

Model Selection: (1) 60% Training, 20% Validation, 20% Test (2) Train different models (3) Check via Validation and select model (4) Use selection on test data!

K-Fold Cross Valid.: not enough data \rightarrow (1) Split 80% data into k parts (2) Round 1 uses first part as **validation** (3) average all accuracy to get final accuracy

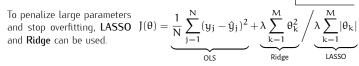
• Underfitting (too loose), Just Right (not perfect fit, but distribution is learnt), Overfitting (too perfect, no distribution)

Regression: X regressor, ε error $y_{real} - y_{lin}$: $Y = X\theta + \varepsilon$ ($y_i = \theta_0 + \theta_1 \cdot x_i + \varepsilon_i$)

Cost Function: used to minimize MSE; $J(\theta_0, \theta_1) = \frac{1}{N} \sum_{i=N}^{N} \epsilon_i^2$

Ordinary Least Squares (OLS): $\frac{\partial J(\theta)}{\partial \theta} = 0 \rightarrow \theta_{opt} = (X^TX)^{-1}X^Ty$ (Gradient **Descent** iteratively: 1 random θ 2 small steps down cost surface \bullet direction = negative gradient; $\theta_{new} = \theta_{old} - \alpha^{\partial J(\theta)/\partial \theta}$

Regression Performance: $R^2 = 1 - \frac{\sum_j \varepsilon_j^2}{\sum_j \varepsilon_\delta^2} = 1 - \frac{\sum_j (y_j - \hat{Y})^2}{\sum_j (y_j - \overline{Y})^2}$ Plot data, as R^2 does not show the distribution (*Anscome's quartet*)!



Polynomial Regression: $h(\theta, X) = \theta_0 + \theta_1 X + \theta_2 X^2 + \cdots \Rightarrow \theta_0 + \theta_1 X_1 + \theta_2 X_2 + \cdots$ Classification =

Classification predicts categories (decision boundry) vs. Regression predicts values (data relationship)

Logistic Regression

$$\hat{Y} = P(Y = 1|X) = h_{\theta}(X) = g(X\theta) = \frac{1}{1 + e^{-X\theta}}$$

$$g(X\theta)$$

$$\hat{Y} = g(X\theta)$$

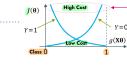
$$g(X\theta)$$

$$g($$

Instead of either exactly 1 or 0, a percentage [0,1] is given. No OLS-form possible, good parameters calculated via Gradient Descent.

Cross Entropy Cost Function

Classifying for one class **must** penalize the other!



One Versus The Rest

Train model based on a specific class, against others. It is either in that class (1) or not $(0) \rightarrow One vs. All classification.$

Performance Analysis =



Dealing with Bias & Variance

Dealing with bias is easier (underfitting): ● larger set of feature ● different features • feature engineering • more complex algorithms (more data is not necessarily the answer)

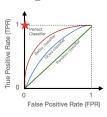
Dealing with variance (overfitting): • reduce features • get more training data • stop training early to avoid overfitting • data cleaning Bias means we are systematically off by a fixed amount. Variance means we are systematically off in a scattered way

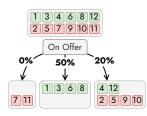
Receiver Operating Characterstic (ROC):

- to compare different models
- largest area under curve is best

Decision Trees —

Gini Impurity measures the likelihood of an incorrect classification of a random data record. *Impure* features produce no significant information. Cont. Var. (1) Sort Table (2) Calculate avg. of adjacent values $(10 \rightarrow 15 \leftarrow 20)$ (3) Averages are splitting criteria (histogramm binning reduces splits)





Label	p(No)	p(Yes)	1-p(No) ² -p(Yes) ²
0%	1	0	0
20%	2/3	1/3	4/9
50%	0	1	0

Gini Impurity of entire feature:

$$2/12 \times 0 + 6/12 \times 4/9 + 4/12 \times 0 = 2/9$$

Regression Trees • Are grown like decision trees for classification but with different splitting criteria • Instead of Gini impurity, minimize variance of values in the same subset • Finally, predict average value of all instances in a leaf node

Tree Construction Rules

(1) **Only** positive or negative nodes left \rightarrow stop branch & assign corresponding decision as leaf node (2) **Some** positive and negative remain \rightarrow apply another feature (3) No instances left \rightarrow look at parent node and decide according the more frequent label (4) Instances left but **no** features \rightarrow training data is no good or contains hidden features; decide on the more frequent label

Pros & Cons Decision & Regression Trees

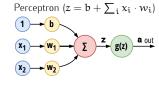
+ readable, can both candle numeric & categorial, (almost) no data preparation, perform well on larger datasets — not as accurate, danger of over-complex trees, very sensitive to data qualitu

Random Forests (collection of decision trees): Utilize random (row)sampling with replacement and random (column) feature selection (without replacement) to be less sensitive & no overfitting.

Ensembles complex ML model built from simpler models. Building Bricks \rightarrow Bragging computes weighted average of votes from simple models, Boosting improves simple model by correcting its errors ($F_m(x) + h_m(x)$, e.g. Gradient Boosting).

Neural Networks

Input x_i from previous 'axons', contain some kind of information / Weights b, w_i define the inputs' influence / Sum \(\sum \) sum of the weighted inputs / Activation q(z) can be hard threshold $(\alpha = g(z) = 1 \text{ if } \sum \geq 0 \text{ else } 0), \text{ logistic}$ regression $(g(z) = \frac{1}{1 + \exp(-z)})$



Feed Forward data moves one way (left → right)

Hidden Layers: allows for new features α to be added

Layer Count $c_L = c_{hidden} + c_{output}$ (input is not counted)

Feed Forward →

- · A neural net is the connection of many artificial neurons in parallel and series
- Each layer of a feed-forward ANN is defined by its weights W and biases b
- Given data with labels **v** and features x, we feed x forward through the ANN

Back Propagation ←

- . At the output of the ANN we get the estimate $\hat{\mathbf{y}}$ of \mathbf{y}
- Because we start with random
- Define the error J as a measure of distance between $\hat{\mathbf{v}}$ and \mathbf{v}
- minimize J. This occurs when the gradient of J is small or zero
- We introduce deltas δ at each

- Back-propagation gives the
- · The gradient is a vector which points steeply up the
- So take a small step α in the opposite direction, reducing J
- "Take a step" means making
- Changes are defined exactly
- · This is gradient descent by feed forward & back-prop

- **W** and **b**, the estimate \hat{y} is poor
- We seek the best W & b to
- So we need the gradient of J
- · Compute the gradient iteratively
- layer to simplify computation

Gradient Descent 1

- gradient of J with respect to weights W and biases b

- small changes to W & b
- by $-\alpha$ times the gradient

Activation Function

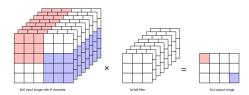
ReLU (rectified linear unit ;good for hidden) max(0, x) brings better perfomance and alleviates the vanishing gradients problem (flat line activation) → introduces **dying units** (bad for gradient training)

Leaky ReLU max(0.1x, x) introduces non-zero for negative values and fixes vanishing gradients + dying unit \rightarrow results aren't always consistent **Soft-Max** (good for output) softmax_j = $\frac{\exp z_j}{\sum_{k=1}^K \exp(z_k)}$ is a generalization of the logistic function to **multiple dimensions** and can be interpreted as



probabilites (normalization). **Convolution Neural Network**

Features are extracted from images using convolution (Faltung). Weights are used as Masks/Filters on these images to extract certain features such as edges or colour. The final layer mostly uses softmax!



Pooling applies operations such as max, min or avg to a pixel neighbour and the neighbourhood size is determined by the pool size parameter.

Parameter Counting Example on 20 \times 20-RGB-Image (1) 2 Filters of 3 \times 3 $((3 \cdot 3 \cdot 3 + 1) \cdot 2 = 56 \text{ results } 18 \times 18 \times 2 \text{ Layers})$ (12) 3 Filters of 3 × 3 with 2 channels each $((3 \cdot 3 \cdot 2 + 1) \cdot 3 = 57 \text{ results } 16 \times 16 \times 3 \text{ Layers})$ (L3) Flatten result to a single column of neurons $(1 \cdot 16 \cdot 16 \cdot 3 = 1 \times 768)$ (L4) Fully connected (aka. dense) layer and 2 output channels (#param = $768 \cdot 2 + 2 = 1538$ (2 biases! & 2 neurons/output shape))

Natural Language Processing

Syntax compare spelling or words / Similarity Levenshtein Distance

Word Relatedness Words are related... • when they appear in the same document

(synonyms not related!!) • when they relate to the same topic (requires a lot of space) when they can be replaced by each other

Word Similarity Words are similar when they appear in the same context

Term Frequency (Inverse Doucment Frequency) TF-IDF(x,y) = TF(x,y) $\cdot \log \left(\frac{N}{DF(x)} \right)$

/ N number of documents in the corpus, TF counts the number of occurences of x in y, DF counts the number of documents that contain x. QUETZALCOATLUS has high score, due to being used a lot in a **single** document, compared to the low scoring word AND (used everywhere).

Recurrent Neural Nets (RNN)

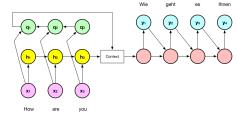
$$h_t = (W_{xh} \cdot x_t + W_{hh} \cdot h_{t-1} + b_h) \quad y_t = \varphi(W_{hy} \cdot h_t + b_y)$$

RNN are made for the supervised analysis of arbitrary time-series data (instead of fixed length, various sized strings can be analyzed) and do not parallelize! LSTM & GRU help with alleviating vanishing gradients to remember more of the context (Training an RNN by multiplying gradients backwards in time).

GRU ($\not\bowtie_{t}$) use hidden states as memory (faster) & Long Short-Term Memory Model have an additional memory state (better performance)

Bidirectional RNN reads left and right \rightarrow produces together output

Encoder-Decoder Architecture: usually RNNs have one input and output, which makes it unusuable for i.e. translators. To allow in- and outputs of various lengths, en- & decoder architecture is used.



Mimics the behaviour of knowing which page the content can be found in. Words are assigned an importance value which is used by the next generation for the output words. In Encoder Attention Decoder Out

k-Means Clustering

(1) Input k Clusters and data points x_n (2) Randomly choose k clusters centers μ_k (3) Repeat until convergence (stability): (a) assign data points to nearest cluster center (b) update cluster center with mean of assigned datapoints

Two data points can meet in a cluster and separate again in the next step! \rightarrow pattern always changes

Clustering Distortion

Total distortion sum of squared distances of points and cluster center • Average **Distorion** average of total distortion (additional 1/n) and allows for comparison with other data sets

k-Means only approximates optimal solutions and converges either to global OR local

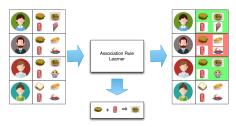
Elbow Methods increasing the amount of clusters leads to an 'elbow' in the distance $nCluster\ diagram \rightarrow use\ cluster\ count\ at\ the\ 'elbow'$

Agglomerative Clustering -

Outputs an hierarchy (dendrogram) that are informative and human interpretable. It does not scale with larger data, but cluster stays together forever! (1) Initially each datapoint is a separate cluster (2) While stop condit. (on no change) is not met: (a) calculate distance between pairs of cluster (b) merge pairs with shortest distance

Association Rules

Implication $(X \to Y)$: the higher confidence, the more likely Y is also present in transaction with X



Support (Interestingness): how frequently an item set occurs in the data $support(\{i_1,\ldots,i_n\}) = \frac{\text{\#purchases of }\{i_1,\ldots,i_n\}}{\text{\#purchases of }\{i_1,\ldots,i_n\}}$

Support of an Association support $(X \to Y) = \text{support}(Y \to X) = \text{support}(X \cup Y)$

Confidence (Trustworthiness): how frequently items in Y appear in transactions that contain X confidence $(X \to Y) = \frac{\text{support}(X \cup Y)}{\text{support}(X)}$

BOTH Support and **Confidence** need to be considered. — Only Antecedent X and co-occurence $X \cup Y$ is taken into account, not Y; sup. & conf. cannot filter coincidental occurrences out.

Lift (Association Strength): how many times more often X and Y show up together than expected lift(X \rightarrow Y) = $\frac{\text{support}(X \cup Y)}{\text{support}(X) \cdot \text{support}(Y)}$ (= 1 independent; < 1 X & Y appear less often together; $> 1 \times 8 \times 9$ appear often together)

Apriori Algorithm

(1) Generate frequent item sets $\rightarrow (1)$ frequent items leads subsets being frequent too (2) infrequent leads supersets infrequent too. (shape lattice of subsets: first layer single node, second double connection,...)

(2) Extract rules from frequent item satisfying the confidence threshold \rightarrow (1) when an item violates the confidence bound all subsets can be eliminated. (shape lattice of

Recommender System - Non-Personalized —

+ simple, fast & cheap; Associations are context aware; Associations can be tailored to specific businesses (who viewed, who bought...); No new user cold start problem non-personalized

Recommender System - Personalized -

$$P_{\mathbf{u}j} = \frac{\sum_{\mathbf{i} \in \mathbf{N}_j} \mathsf{similar}(j, \mathbf{i}) \cdot r_{\mathbf{u}\mathbf{i}}}{\sum_{\mathbf{i} \in \mathbf{N}_i} \mathsf{similar}(j, \mathbf{i})} \quad P_{\mathbf{u}j} = \overline{r}_{\mathbf{u}} + \frac{\sum_{\mathbf{i} \in \mathbf{N}_\mathbf{u}} \mathsf{similar}(\mathbf{u}, \mathbf{i}) \cdot (r_{\mathbf{i}j} - \overline{r}_{\mathbf{i}})}{\sum_{\mathbf{i} \in \mathbf{N}_\mathbf{u}} \mathsf{similar}(\mathbf{u}, \mathbf{i})}$$

N rated items (hyperparameter); P_{uj} how much does user u like item j; N_i Neighbours of item i; r_{ui} rating that user u gave item i; N_u Neighbors of user u; r_{ij} from neighbor i to item j

Personalized Content-Based Recommendations (left equ.): + user independence, no new item cold start problem — limited content analysis, overspecialization (bubble esque), severe new user cold start problem

User-to-User Collaborative Filtering (right equ.):

+ cross-category recommendations many participants required, system must first learn user + item rating (cold start problem), gray sheep problem (unusual taste results in low accuracy)

Hybridization

Combination of different recommender systems! • Weighted Approach item score is weighted sum score from different recommenders • Mixed Approach results of different recommender are presented • Cascade Approach one recommender refines the result of other method • Switching Approach use different methods ta different places (online