



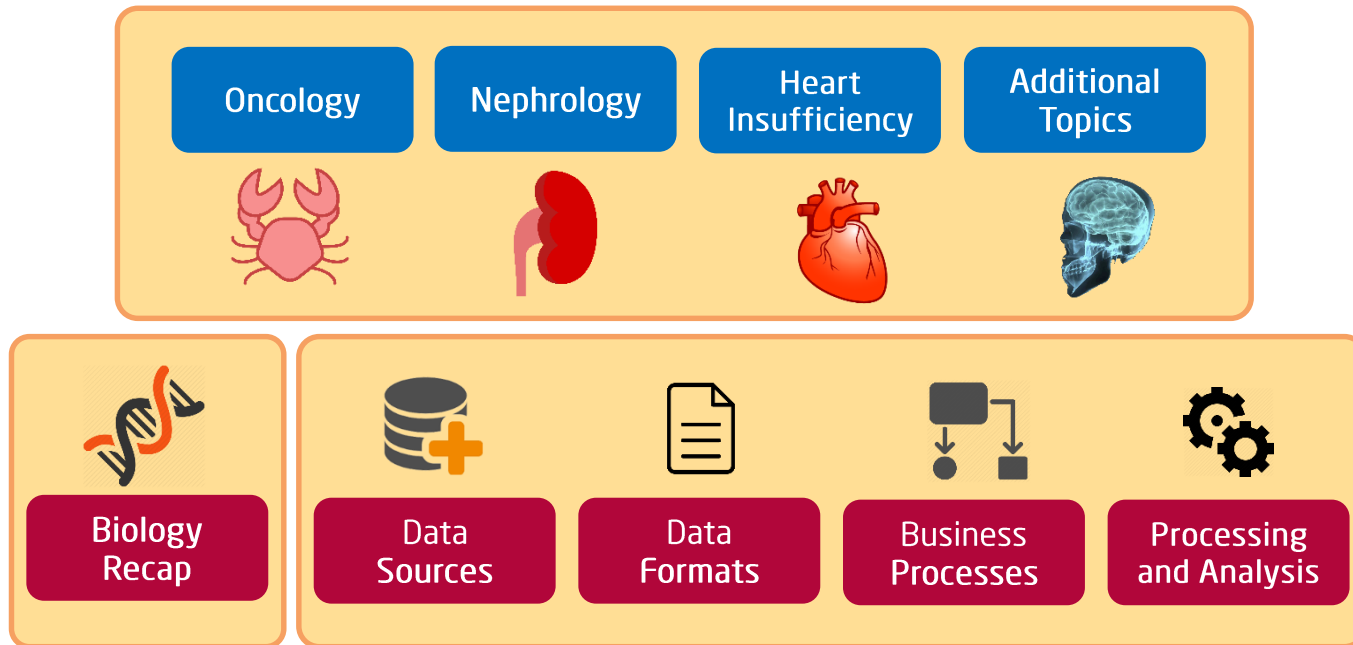
# Developing Clinical Prediction Models for Nephrology

Harry Freitas da Cruz  
Data Management for Digital Health  
Summer 2017

# Where are we?

Data Management  
& Foundations

Real-world  
Use Cases



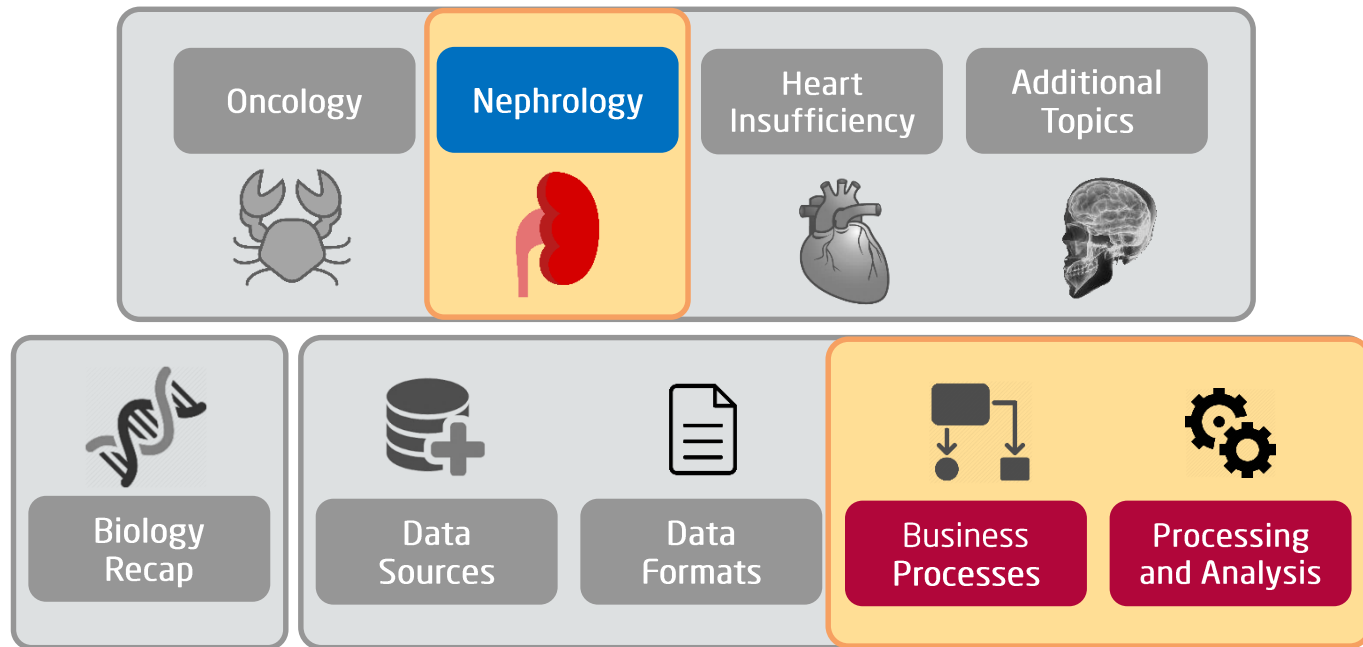
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# Where are we?

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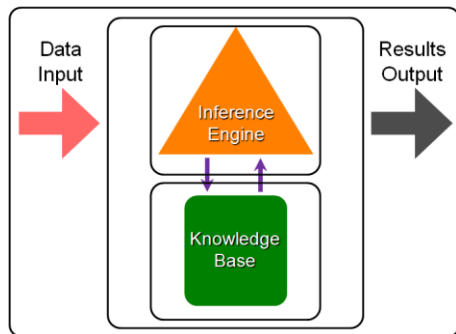
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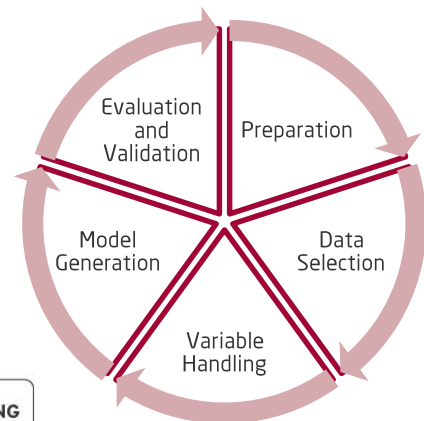
# Recap Predictive Modelling for Clinical Applications

- Predictive analytics in healthcare
- Clinical Decision Support Systems
- Clinical Data Repository
- Developing a Clinical Prediction Model

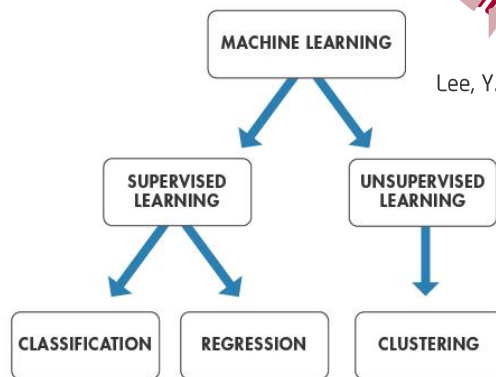
- Preparation
- Data selection
- Variable handling
- Model generation



Architecture components of CDSS (Kola, n.d.)



Lee, Y.-H., Bang, H., & Kim, D. J. (2016)



<https://de.mathworks.com/help/stats/machine-learning-in-matlab.html>

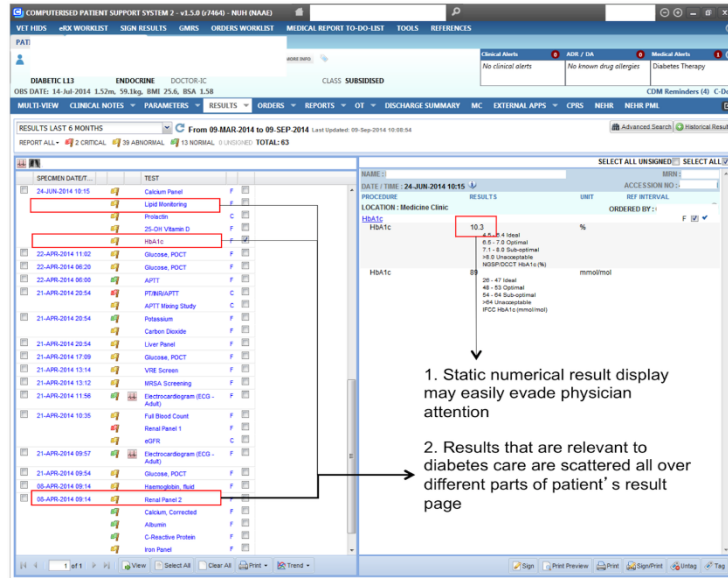
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# Clinical Decision Support Systems

## More Examples

### ■ Diabetes Dashboard<sup>1</sup>



A screenshot of Computerized Patient Support System, Sim (2017)



A screenshot of the summary dashboard, Sim (2017)

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# Clinical Decision Support Systems

## More Examples

### ■ National Kidney Foundation (US)



EGFR Calculator (NKF)



iPhone App (NKF)



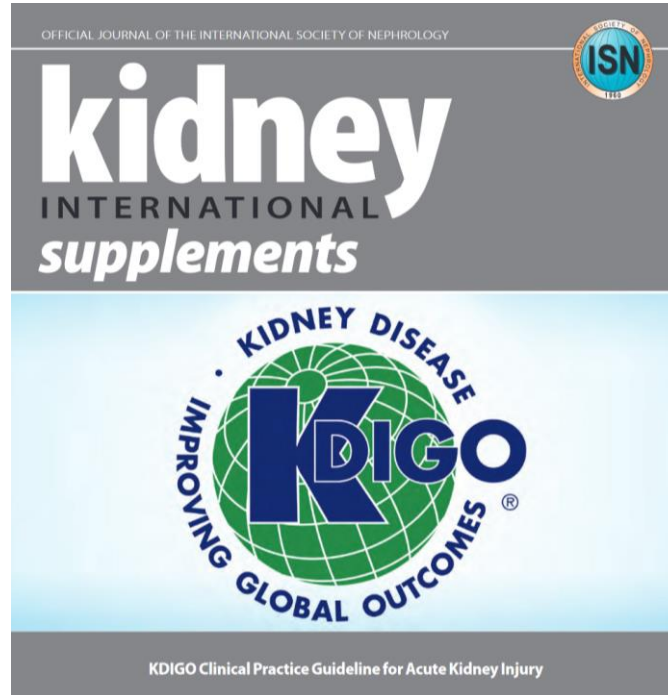
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# Clinical Guidelines

## KDIGO – Kidney Disease Improving Global Outcomes



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# Clinical Guidelines

## KDIGO – Kidney Disease Improving Global Outcomes

### ■ Recommendation level

Grade*	Implications		
	Patients	Clinicians	Policy
<b>Level 1</b> “We recommend”	Most people in your situation would want the recommended course of action and only a small proportion would not.	Most patients should receive the recommended course of action.	The recommendation can be evaluated as a candidate for developing a policy or a performance measure.
<b>Level 2</b> “We suggest”	The majority of people in your situation would want the recommended course of action, but many would not.	Different choices will be appropriate for different patients. Each patient needs help to arrive at a management decision consistent with her or his values and preferences.	The recommendation is likely to require substantial debate and involvement of stakeholders before policy can be determined.

### ■ Quality of evidence

Grade	Quality of evidence	Meaning
<b>A</b>	High	We are confident that the true effect lies close to that of the estimate of the effect.
<b>B</b>	Moderate	The true effect is likely to be close to the estimate of the effect, but there is a possibility that it is substantially different.
<b>C</b>	Low	The true effect may be substantially different from the estimate of the effect.
<b>D</b>	Very low	The estimate of effect is very uncertain, and often will be far from the truth.

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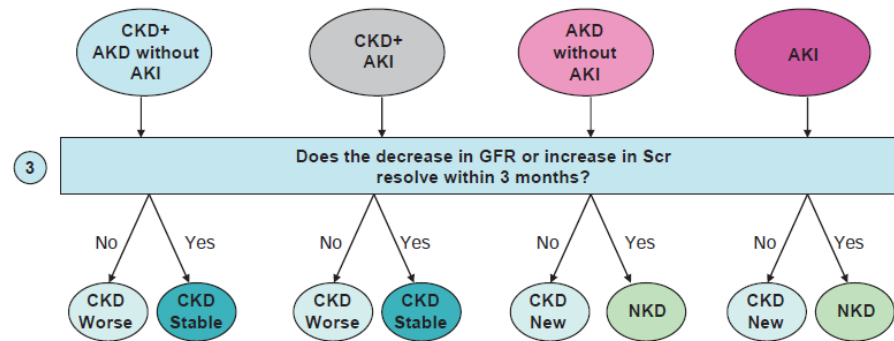


# Clinical Guidelines

## KDIGO – Kidney Disease Improving Global Outcomes

### ■ Recommendations and flowcharts

- 2.2.1:** We recommend that patients be stratified for risk of AKI according to their susceptibilities and exposures. *(1B)*
- 2.2.2:** Manage patients according to their susceptibilities and exposures to reduce the risk of AKI (see relevant guideline sections). *(Not Graded)*
- 2.2.3:** Test patients at increased risk for AKI with measurements of SCr and urine output to detect AKI. *(Not Graded)* Individualize frequency and duration of monitoring based on patient risk and clinical course. *(Not Graded)*



KDIGO Guidelines, GFR/SCr algorithm: excerpt

KDIGO Guidelines, Risk assessment of AKI

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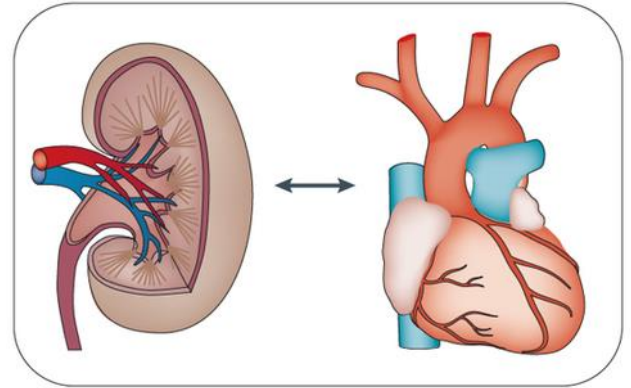
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- Developing a Clinical Prediction Model for predicting AKI in heart surgery
- Tools for developing a Clinical Prediction Model
- Validating and evaluating models (ROC Curve)
- Use Case Nephrology exercise: what to expect?

# Developing a Clinical Prediction Model Incidence of Acute Kidney Injury in Cardiac Surgery

- Heart and kidney are deeply connected
- AKI after cardiac surgery is relatively common (3 to 30%)
- Associated with complications and mortality
- Risk factors: age, hypertension, vascular disease (i.a.)
- Cardiopulmonary bypass (CPB)
- Body fluid volume alterations
- High doses of vasopressors



<http://www.nature.com/nrneph/journal/v12/n10/abs/nrneph.2016.113.html>

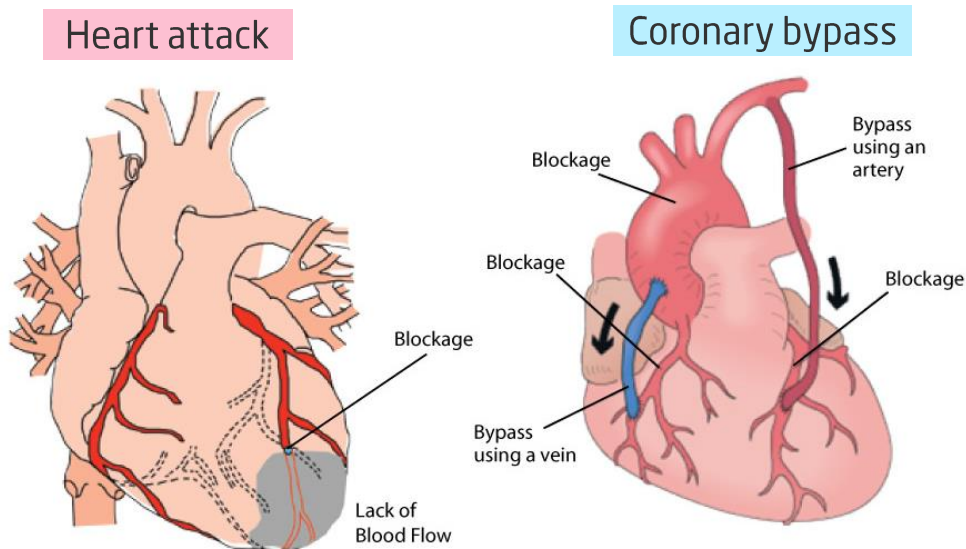
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# Developing a Clinical Prediction Model Bypass and Valve Surgery

- Coronary artery bypass graft (CABG)
- ICD-9-PCS: Ch. 36.1 - Bypass anastomosis for heart revascularization



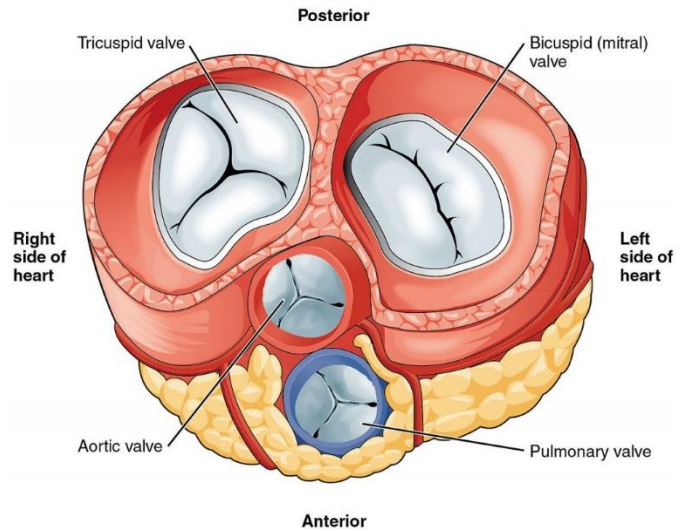
<http://ctsurgery.com/heart-surgery/>

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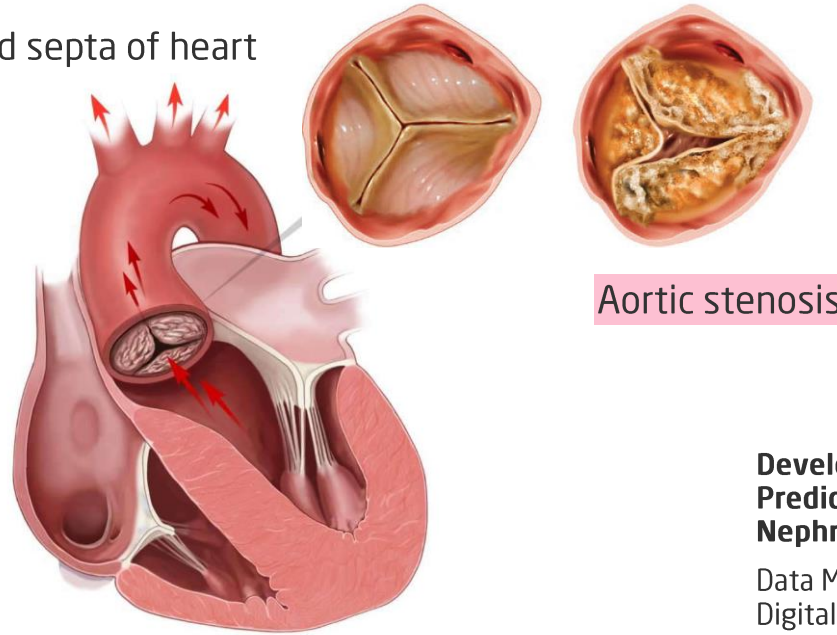
# Developing a Clinical Prediction Model Bypass and Valve Surgery

- Valve surgery
- ICD-9-PCS: Ch. 35 - Operations on valves and septa of heart



<http://teachmeanatomy.info/thorax/organs/heart/heart-valves/>

Normal aortic valve



<http://www.corevalve.com/us/what-severe-aortic-stenosis/index.htm>

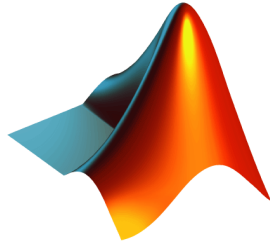
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# Developing a Clinical Prediction Model

## Tools for Developing CPM



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# Developing a Clinical Prediction Model

## Tools: tranSMART

- Clinical and omics data repository
- Initial development at J&J
- Open-source project at the tranSMART Foundation
- Platform for biomedical computation
  - Clinical trials + omics
- Types of analyses (excerpt)
  - Correlation / regression
  - K-means clustering,
  - Survival analysis, aCGH, GWAS



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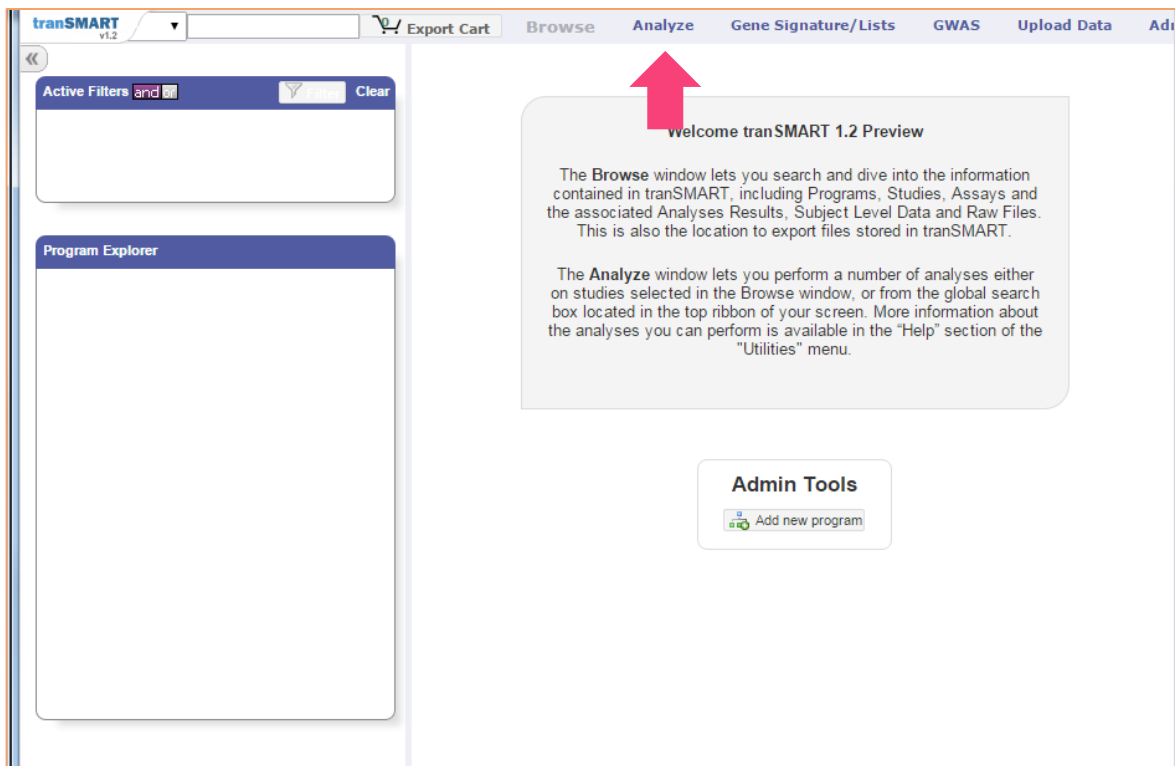
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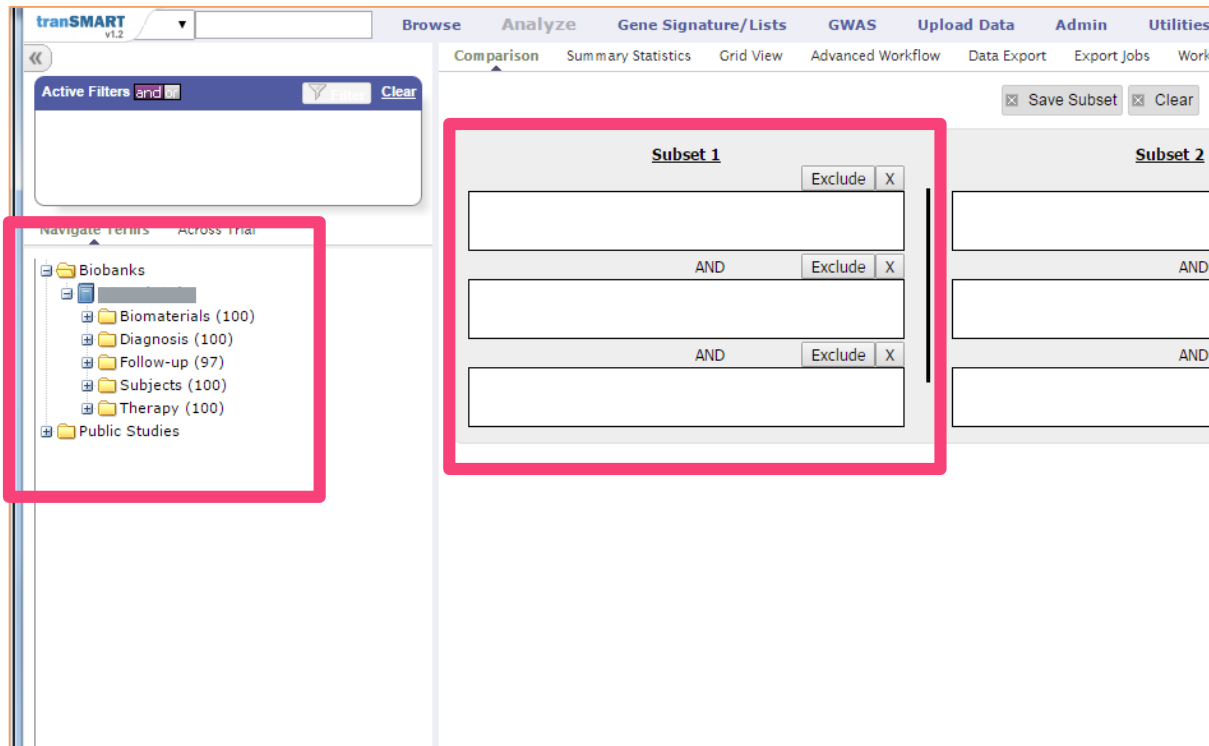
# Developing a Clinical Prediction Model

## Tools: tranSMART



# Developing a Clinical Prediction Model

## Tools: tranSMART

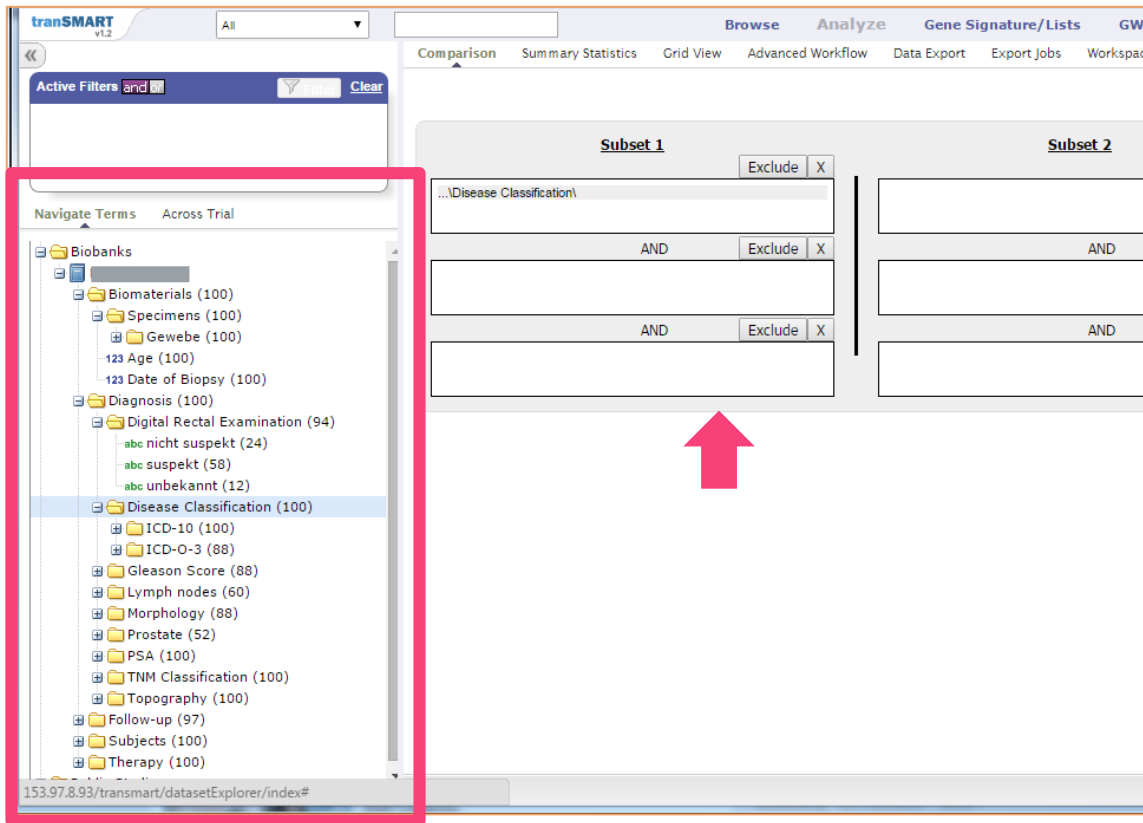


The screenshot displays the tranSMART v1.2 web interface. The top navigation bar includes tabs for Browse, Analyze (selected), Gene Signature/Lists, GWAS, Upload Data, Admin, and Utilities. Below this, a secondary navigation bar shows options like Comparison, Summary Statistics, Grid View, Advanced Workflow, Data Export, Export Jobs, and Work. The main interface is divided into several sections:

- Active Filters:** Located at the top left, it shows a filter set named 'and' with a 'Clear' button.
- Navigate Terms:** A sidebar on the left lists various categories and their counts: Biobanks, Biomaterials (100), Diagnosis (100), Follow-up (97), Subjects (100), Therapy (100), and Public Studies. This section is highlighted with a red box.
- Subset 1:** A central configuration area for defining a subset. It contains three rows, each with a text input field, a logical operator (AND), and an 'Exclude' button. The first row is also labeled 'Subset 1' and has an 'X' button. This section is highlighted with a red box.
- Subset 2:** A similar configuration area on the right, currently empty, labeled 'Subset 2'.
- Buttons:** At the top right, there are buttons for 'Save Subset' and 'Clear'.

# Developing a Clinical Prediction Model

## Tools: tranSMART



The screenshot displays the tranSMART v1.2 web interface. On the left, a hierarchical tree under 'Navigate Terms Across Trial' lists various clinical data categories and their counts. The 'Disease Classification' category is highlighted. On the right, a comparison table is shown with two columns, 'Subset 1' and 'Subset 2', and rows for different clinical terms. A red arrow points to the 'Disease Classification' row in the 'Subset 1' column.

**tranSMART v1.2**

Active Filters: and [X] Clear

**Navigate Terms Across Trial**

- Biobanks
  - Biomaterials (100)
  - Specimens (100)
    - Gewebe (100)
  - Age (100)
  - Date of Biopsy (100)
- Diagnosis (100)
  - Digital Rectal Examination (94)
    - abc nicht suspekt (24)
    - abc suspekt (58)
    - abc unbekannt (12)
  - Disease Classification (100)**
    - ICD-10 (100)
    - ICD-O-3 (88)
    - Gleason Score (88)
    - Lymph nodes (60)
    - Morphology (88)
    - Prostate (52)
    - PSA (100)
    - TNM Classification (100)
    - Topography (100)
  - Follow-up (97)
  - Subjects (100)
  - Therapy (100)

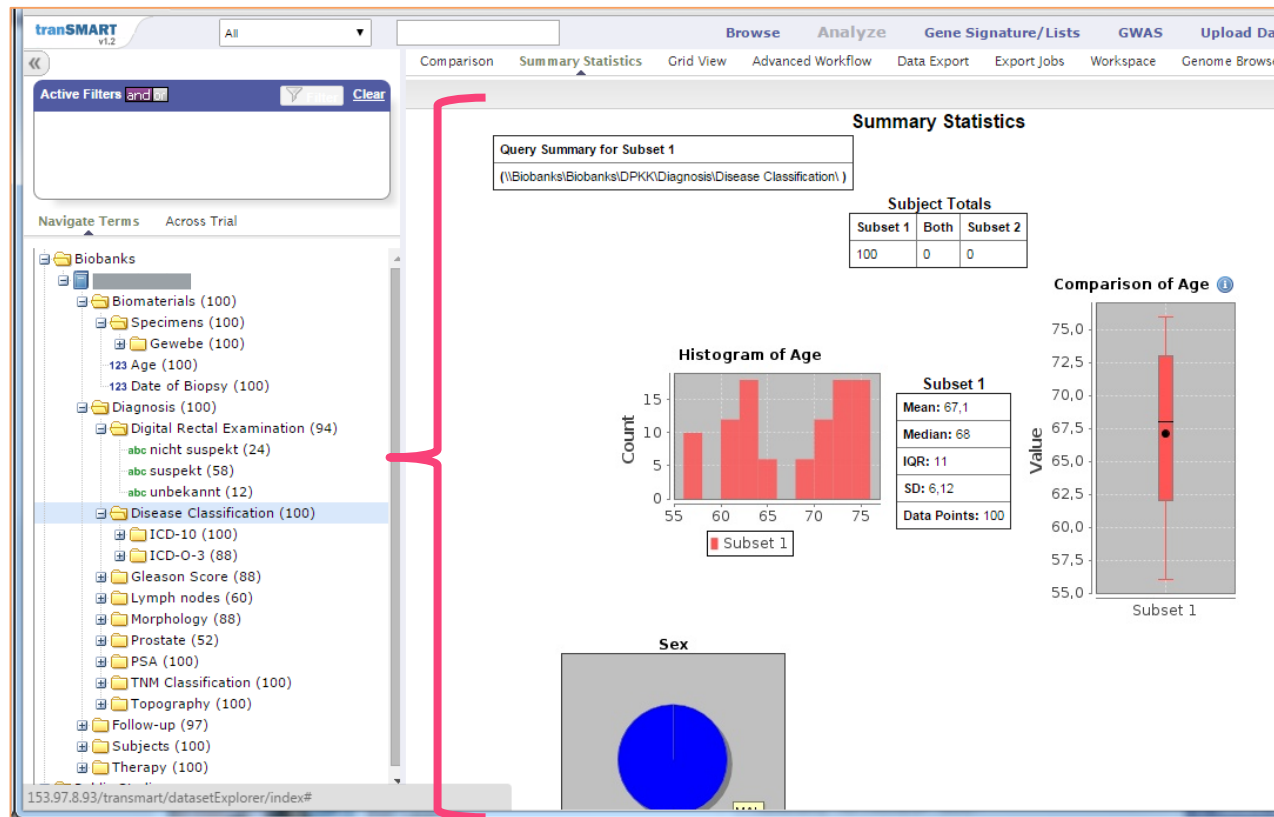
**Comparison**

Subset 1	Subset 2
...Disease Classification	
AND	AND
AND	AND

153.97.8.93/transmart/datasetExplorer/index#

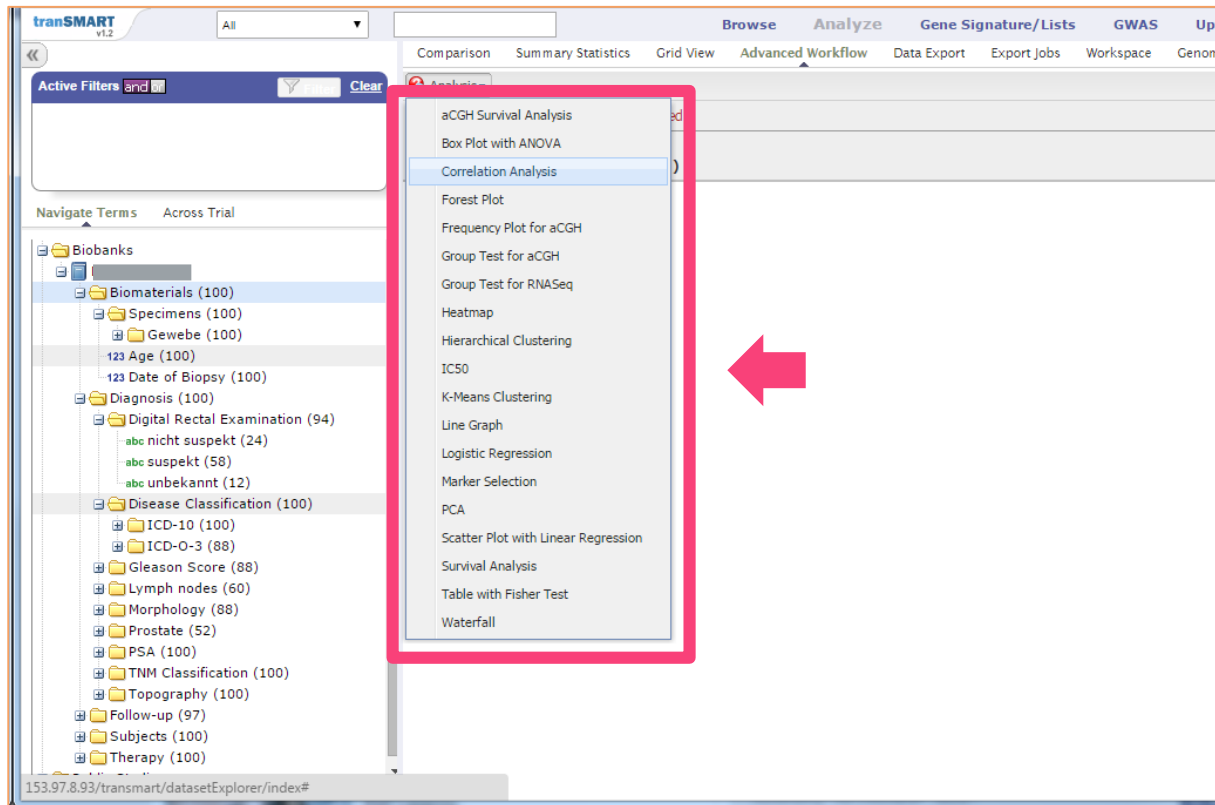
# Developing a Clinical Prediction Model

## Tools: tranSMART



# Developing a Clinical Prediction Model

## Tools: tranSMART



The screenshot displays the tranSMART v1.2 web interface. On the left, a tree view under 'Biobanks' shows a hierarchy of data categories: Biomaterials (100), Specimens (100), Gewebe (100), 123 Age (100), 123 Date of Biopsy (100), Diagnosis (100), Digital Rectal Examination (94) (with sub-items: abc nicht suspekt (24), abc suspekt (58), abc unbekannt (12)), Disease Classification (100) (with sub-items: ICD-10 (100), ICD-O-3 (88), Gleason Score (88), Lymph nodes (60), Morphology (88), Prostate (52), PSA (100), TNM Classification (100), Topography (100)), Follow-up (97), Subjects (100), and Therapy (100). The 'Analyze' menu is open, showing a list of statistical and data analysis tools. A pink arrow points to the 'Correlation Analysis' option in this menu.

tranSMART v1.2

Active Filters and [icon] Clear

Navigate Terms Across Trial

Biobanks

- Biomaterials (100)
  - Specimens (100)
  - Gewebe (100)
  - 123 Age (100)
  - 123 Date of Biopsy (100)
- Diagnosis (100)
  - Digital Rectal Examination (94)
    - abc nicht suspekt (24)
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    - abc unbekannt (12)
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  - ICD-O-3 (88)
  - Gleason Score (88)
  - Lymph nodes (60)
  - Morphology (88)
  - Prostate (52)
  - PSA (100)
  - TNM Classification (100)
  - Topography (100)
- Follow-up (97)
- Subjects (100)
- Therapy (100)

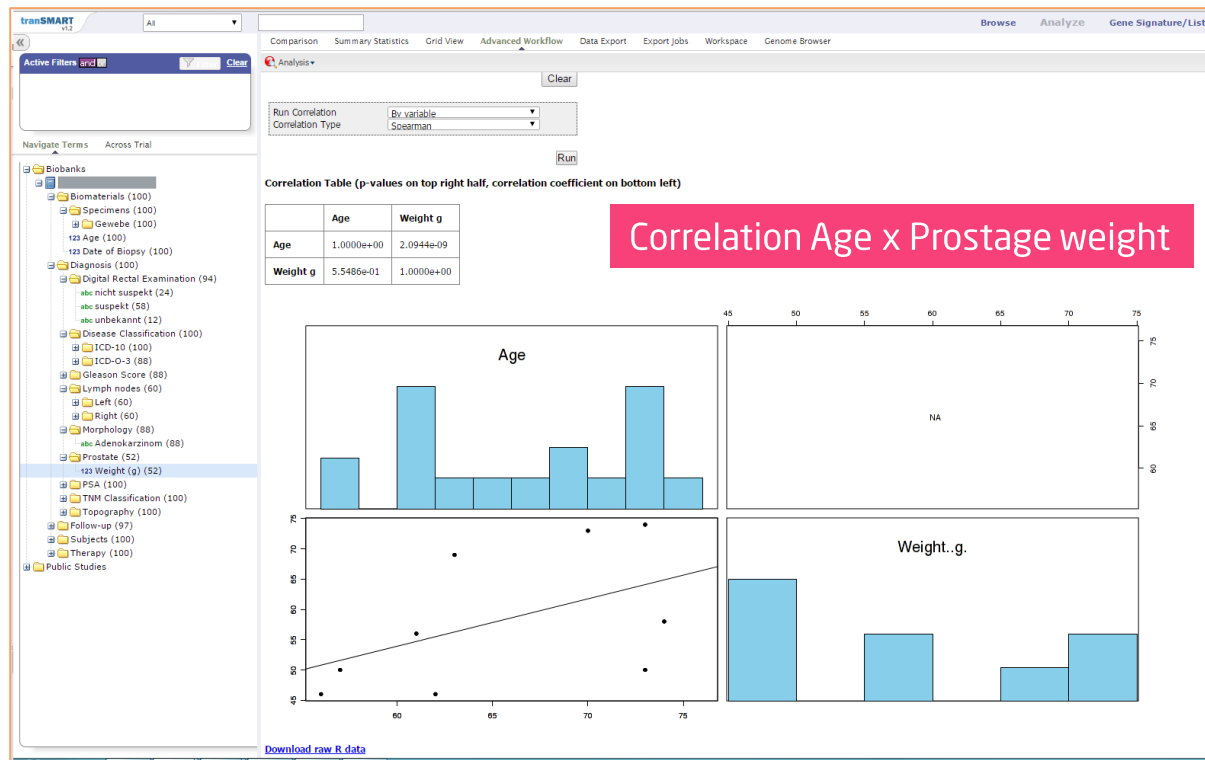
Analysis

- aCGH Survival Analysis
- Box Plot with ANOVA
- Correlation Analysis
- Forest Plot
- Frequency Plot for aCGH
- Group Test for aCGH
- Group Test for RNASeq
- Heatmap
- Hierarchical Clustering
- IC50
- K-Means Clustering
- Line Graph
- Logistic Regression
- Marker Selection
- PCA
- Scatter Plot with Linear Regression
- Survival Analysis
- Table with Fisher Test
- Waterfall

153.97.8.93/transmart/datasetExplorer/index#

# Developing a Clinical Prediction Model

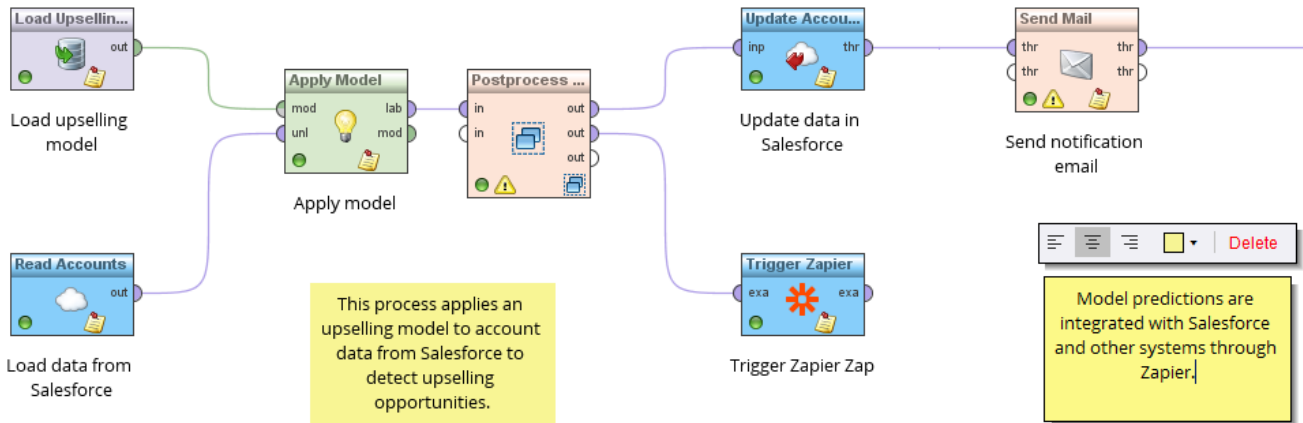
## Tools: tranSMART



# Developing a Clinical Prediction Model

## Tools: RapidMiner

- General purpose ML-prototyping tool
- Process-based (operators)
- Data acquisition, preparation, model development and validation

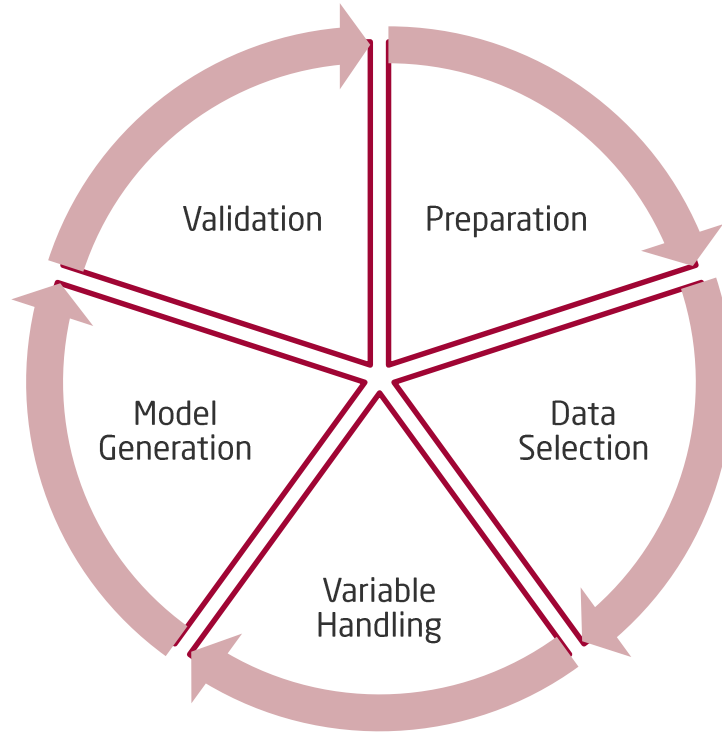


<https://docs.rapidminer.com/studio/releases/changes-6.4.000.html>



# Developing a Clinical Prediction Model

## General approach

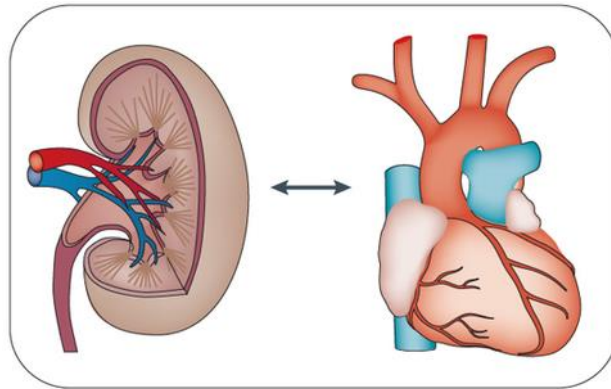


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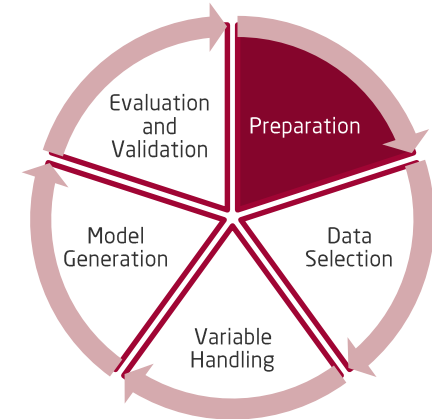
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# Developing a Clinical Prediction Model Preparation

- Target outcome := predict AKI in patients who had cardiac surgery
- Target patients := patients who had cardiac surgery with AKI
- Target users := cardiologists (expert users)



<http://www.nature.com/hrneph/journal/v12/n10/abs/nrneph.2016.113.html>



Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

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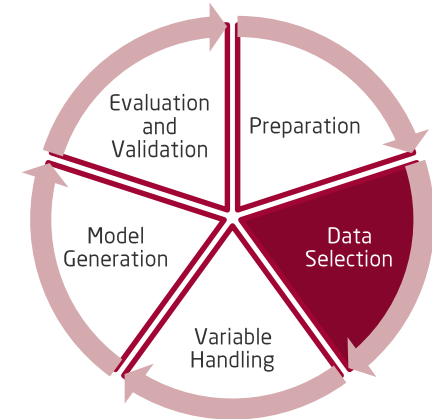
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# Developing a Clinical Prediction Model

## Data Selection

- Patients who underwent *cardiac surgery*
- Patients who presented *AKI*
- Available data source := MIMICIII database<sup>1</sup>

Procedure / Diagnosis	Samples
Cardiac surgery	13,083
Bypass surgery	9,274
Valve surgery	3,809
Acute Kidney Injury (AKI)	1,262
Prevalence	9,65%



Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

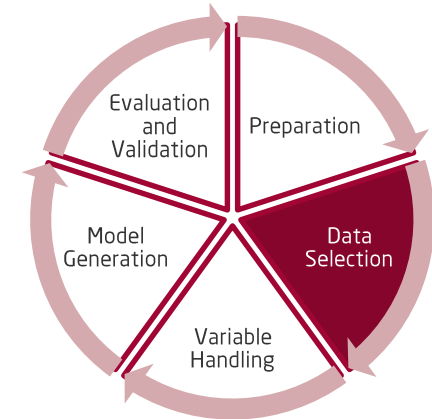
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# Developing a Clinical Prediction Model

## Data Selection

- Data extracted
  - Demographics: gender, age, ethnicity
  - Lab values (min, max): albumin, creatinine, urea, etc.
  - Comorbidities: diabetes, hypertension, obesity, etc.
  - ICU scores: SOFA, SAPS, OASIS
- In total: 78 attributes



Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

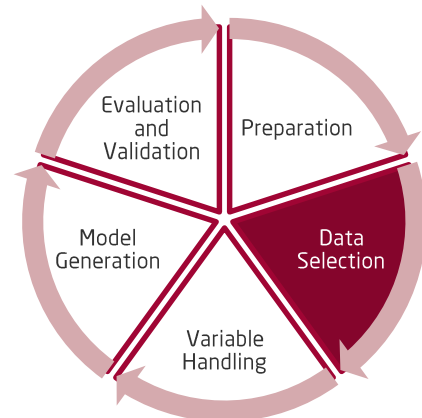
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# Developing a Clinical Prediction Model

## Data Selection: SQL is your friend!

```
select pat.subject_id, adm.hadm_id, pat.gender, adm.ethnicity, round(days_between(pat.dob
, adm.admstarttime)/365,0) as age, elix_score.elixhauser_vanwalraven, aids, alcohol_abuse,
blood_loss_anemia, cardiac_arrhythmias, chronic_pulmonary, coagulopathy,
congestive_heart_failure, deficiency_anemias, depression, diabetes_complicated,
diabetes_uncomplicated, drug_abuse, fluid_electrolyte, hypertension, hypothyroidism,
liver_disease, lymphoma, metastatic_cancer, obesity, other_neurological, paralysis,
peptic_ulcer, peripheral_vascular, psychoses, pulmonary_circulation, renal_failure,
rheumatoid_arthritis, solid_tumor, valvular_disease, weight_loss, sofa.sofa as sofa_total,
sofa.renal as sofa_renal, saps.saps as saps_total, oasis.oasis as oasis_total,
albumin_max, albumin_min, aniongap_max, aniongap_min, bands_max, bands_min,
bicarbonate_max, bicarbonate_min, bilirubin_max, bilirubin_min, bun_max, bun_min,
chloride_max, chloride_min, creatinine_max, creatinine_min, glucose_max,
glucose_min, hematocrit_max, hematocrit_min, hemoglobin_max, hemoglobin_min,
inr_max, inr_min, lactate_max, lactate_min, platelet_max, platelet_min,
potassium_max, potassium_min, pt_max, pt_min, ptt_max, ptt_min, sodium_max,
sodium_min, wbc_max, wbc_min, hsa.aki, case when (hsa.icd_proc between 3610 and
3619) then 'Bypass surgery' when (hsa.icd_proc between 3500 and 3599) then 'Valve surgery'
end from mimiciii.admissions adm inner join mimiciii.patients pat on adm.
subject_id = pat.subject_id inner join mimiciii.m_elixhauser_quan elix on elix.
hadm_id = adm.hadm_id inner join mimiciii.m_elixhauser_quan_score elix_score on
elix_score.hadm_id = adm.hadm_id inner join (select * from (select
sofa.*, min(icustay_id) over(partition by subject_id, hadm_id) as min_icustay_id
from mimiciii.m_sofa sofa ) where icustay_id = min_icustay_id ) sofa on sofa.hadm_id
= adm.hadm_id inner join (select * from (select saps.*, min(
icustay_id) over(partition by subject_id, hadm_id) as min_icustay_id from mimiciii.
m_saps saps ) where icustay_id = min_icustay_id ) saps on saps.hadm_id = adm.hadm_id
inner join (select * from (select oasis.*, min(icustay_id)
over(partition by subject_id, hadm_id) as min_icustay_id from mimiciii.m_oasis oasis
) where icustay_id = min_icustay_id ) oasis on oasis.hadm_id = adm.hadm_id inner join
(select * from (select labs.*, min(icustay_id) over(partition by
subject_id, hadm_id) as min_icustay_id from mimiciii.m_labsfirstday labs )
where icustay_id = min_icustay_id ) labs on labs.hadm_id = adm.hadm_id inner join
mimiciii.heart_surgery_aki hsa on hsa.hadm_id = adm.hadm_id order by pat.subject_id, adm.
hadm_id
```



Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

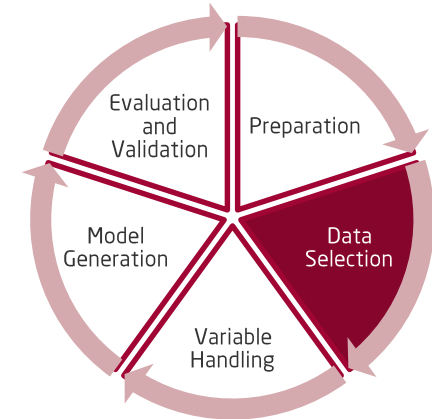
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# Developing a Clinical Prediction Model

## Data Selection

- Data extracted
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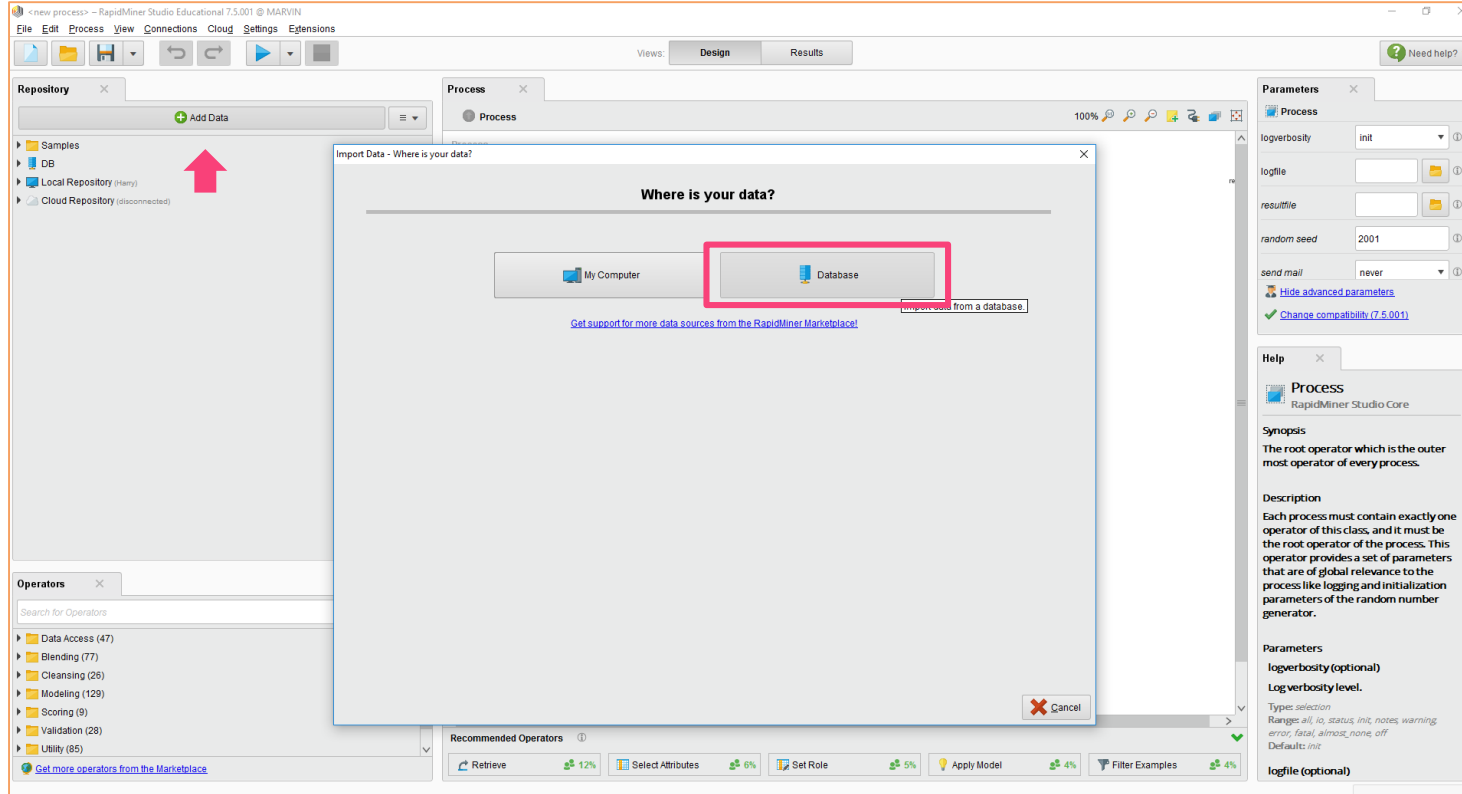
Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

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# Developing a Clinical Prediction Model

## Data Selection: Loading Data

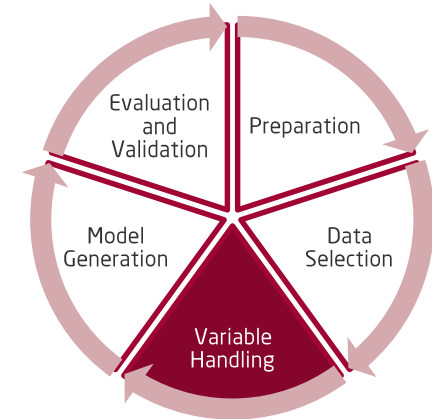




# Developing a Clinical Prediction Model

## Variable Handling

- Watch out for non-relevant predictors (e.g. IDs, etc.)
- Some variables might not be related at all with the output
- Inspect missing values: consider removing variables
- Are any data transformations needed?
  - Highly dependent on the underlying algorithm
  - Normalization and log scale
  - Nominal to numerical



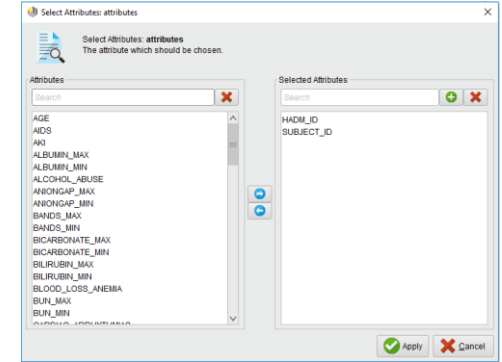
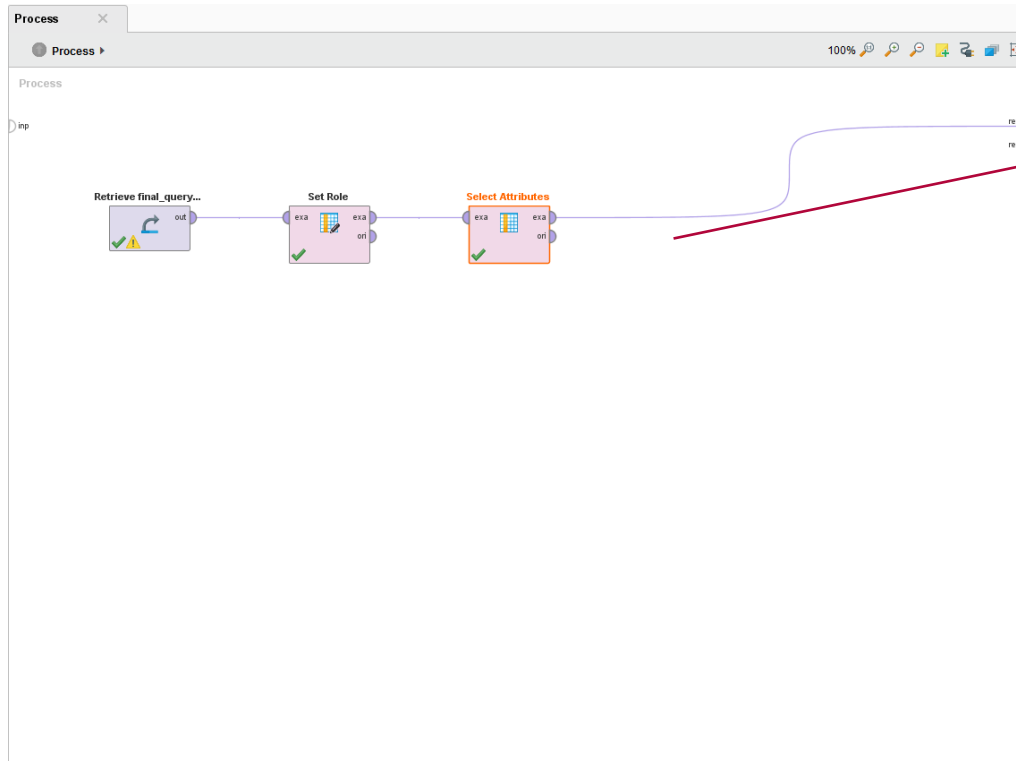
Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

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# Developing a Clinical Prediction Model

## Data Selection and Variable Handling



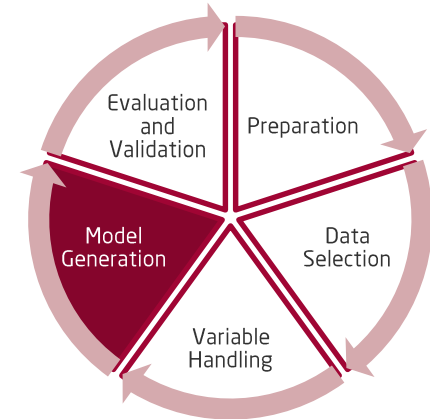
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# Developing a Clinical Prediction Model

## Model Generation

- Start with simple models (Occam's razor)
- Try out some algorithms and compare performance
- One of the most used in medicine: decision trees
- Consider performing feature selection
- Algorithm parameter tuning



Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

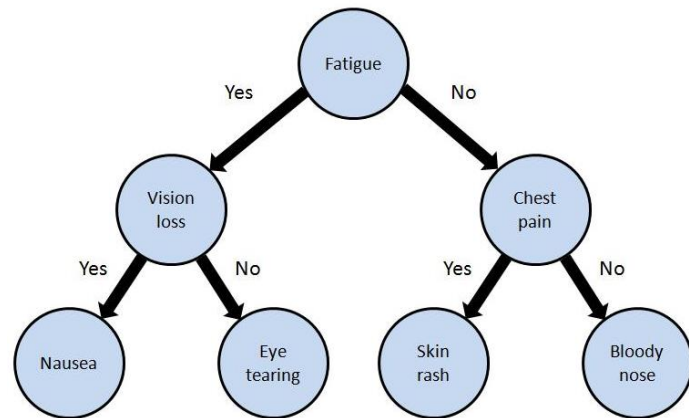
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# Developing a Clinical Prediction Model

## Model Generation: Decision Trees

- Decision rules inferred from data
- Advantages:
  - Interpretability („white box“)
  - Can often be combined with other algorithms
  - Requires little data preparation
- Disadvantages
  - As dimensions increase, so does complexity
  - May lack generalization, prone to overfitting
  - Creates bias if classes are unbalanced



[http://web.eecs.umich.edu/~cscott/research/decision\\_tree.jpg](http://web.eecs.umich.edu/~cscott/research/decision_tree.jpg)

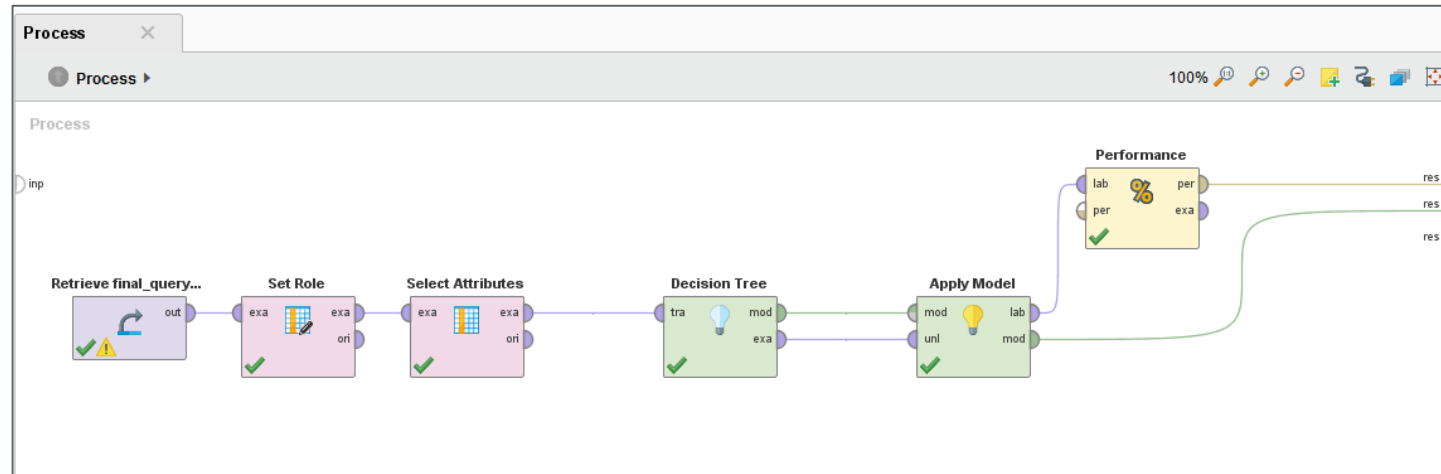
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# Developing a Clinical Prediction Model

## Model Generation: RapidMiner

- Decision tree with no validation



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# Developing a Clinical Prediction Model

## Model Generation: RapidMiner

- Model performance: precision and recall?

Result History x PerformanceVector (Performance) x ExampleSet (//Local Repository/data/final\_query\_v12) x

Criterion

- accuracy
- precision
- recall
- AUC (optimistic)
- AUC
- AUC (pessimistic)

Performance

Description

Annotations

☒ Table View ☐ Plot View

accuracy: 91.81%

	true no	true yes	class precision
pred. no	11787	1037	91.91%
pred. yes	34	225	86.87%
class recall	99.71%	17.83%	

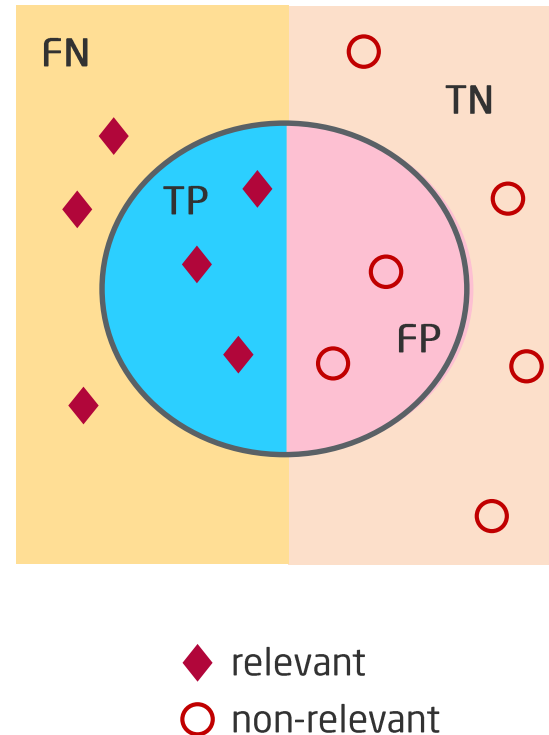
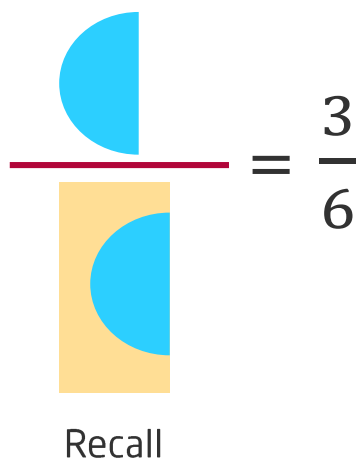
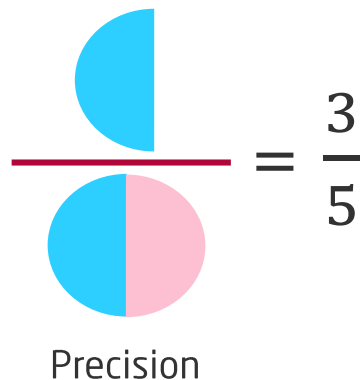
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# Clinical Prediction Models

## Precision and Recall

- Both are relevance measures
- Precision := % retrieved instances that are relevant
- Recall := % relevant documents in the result set



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# Clinical Prediction Models

## Confusion Matrix

- Precision :=  $\frac{TP}{TP + FP}$

- Recall :=  $\frac{TP}{TP + FN}$

- F-measure :=  $2 \cdot \frac{Precision \cdot Recall}{Precision + Recall}$

- Sensitivity := True Positive Rate := Recall

- Specificity := True Negative Rate

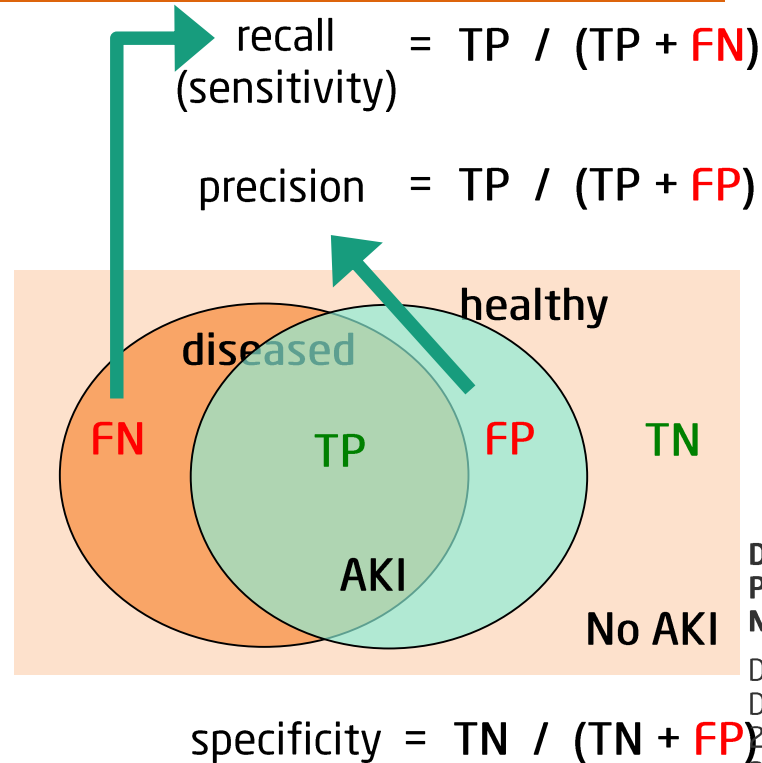
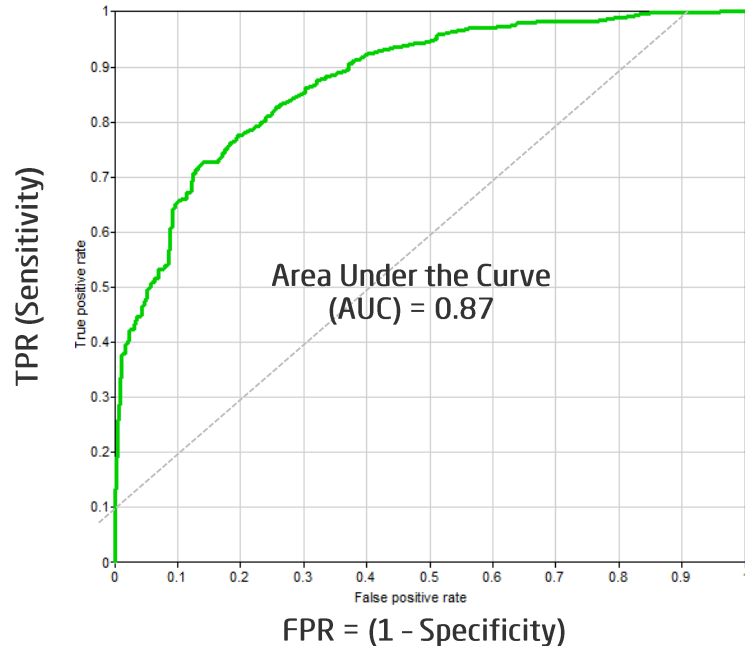
		Predicted	
		Positive	Negative
Actual	Positive	True Positive	False Negative
	Negative	False Positive	True Negative

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# AKI prediction (classifier accuracy)

## Receiver Operating Characteristic (ROC) curve



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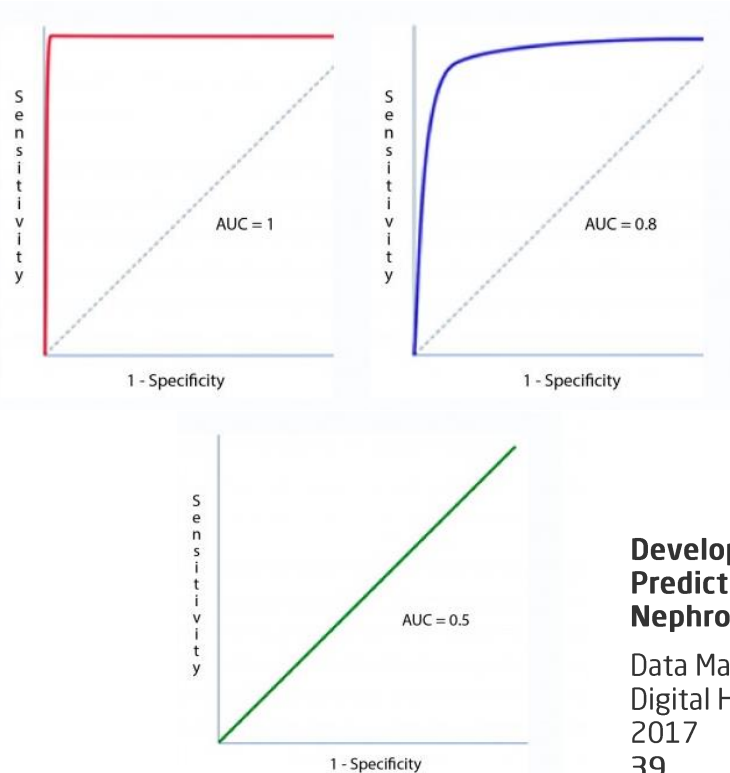
# Clinical Prediction Models

## Receiver Operating Characteristic (ROC) Curve

- Performance of a binary classifier
- Plot showing TPR and FPR
- Varying *classification thresholds*
- Allows comparison between classifiers
- The higher the AUC (Area Under the Curve) the better

$$TPR = \frac{TP}{\underbrace{TP + FN}_{\text{all positive instances}}}$$

$$FPR = \frac{FP}{\underbrace{FP + TN}_{\text{all negative instances}}}$$



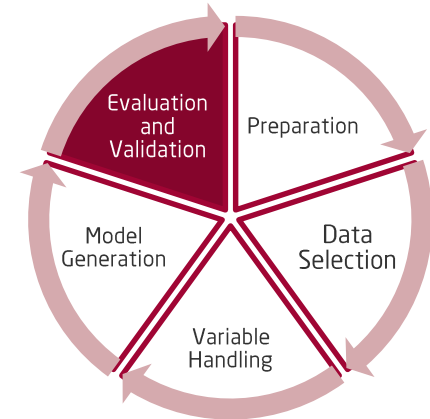
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# Developing a Clinical Prediction Model

## Evaluation and Validation

- Apply cross-validation to assess generalizability
- Try out different algorithms
- Assess the impact of missing values
  - Complete case analysis
  - Single or multiple imputation
- Assess whether class data is balanced
- Perform feature selection



Lee, Y.-H., Bang, H., & Kim, D. J. (2016)

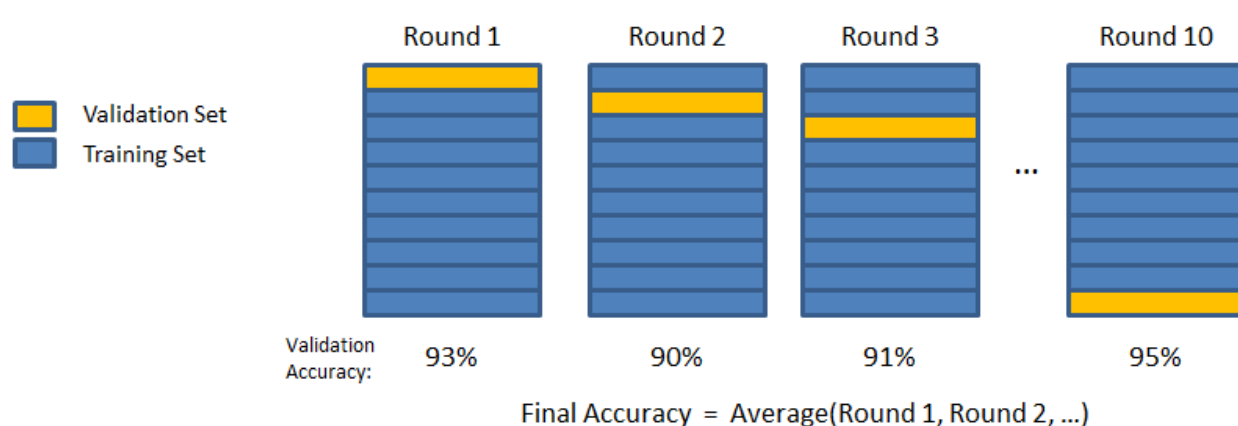
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# Developing a Clinical Prediction Model

## Cross-validation ( $k$ -fold)

- How general is the model?
- Important to avoid *overfitting*
- Useful for internal validation of a model
- Training and validation sets
- Random split into  $k$  subsamples



<https://chrismccormick.files.wordpress.com/>

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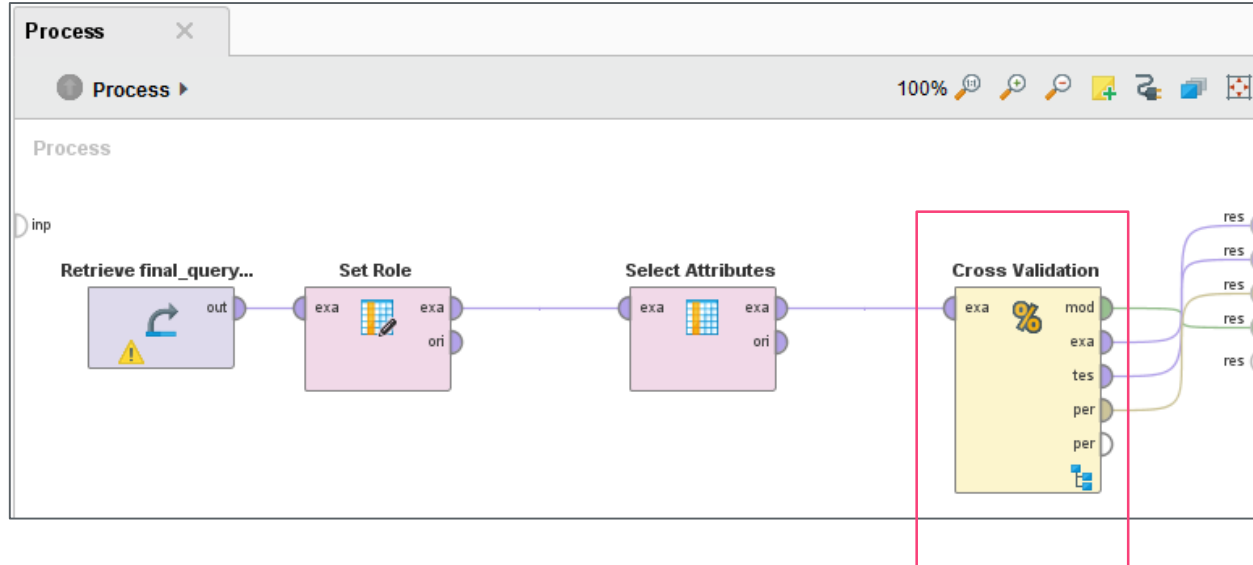
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# Developing a Clinical Prediction Model

## Evaluation and Validation: RapidMiner

- Cross validation operator



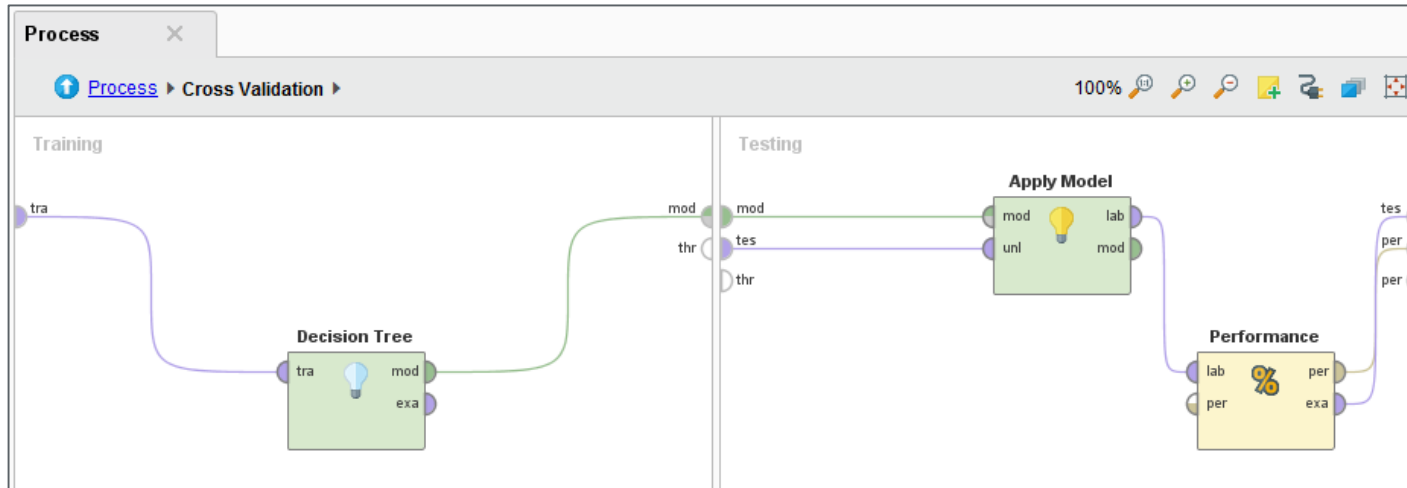
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# Developing a Clinical Prediction Model

## Evaluation and Validation: RapidMiner

- Cross-validation operator



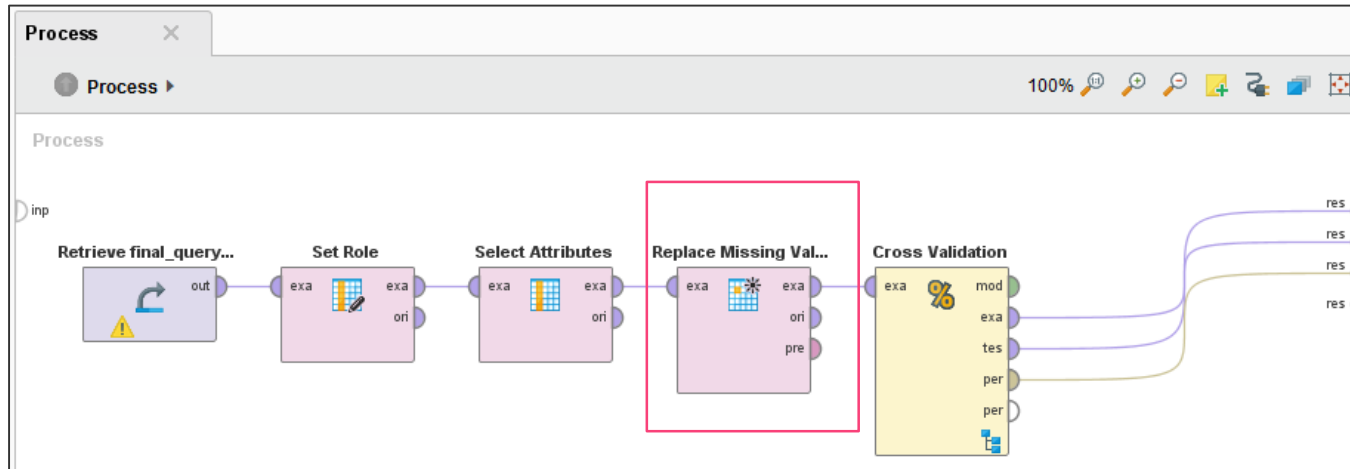
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# Developing a Clinical Prediction Model

## Evaluation and Validation: RapidMiner

- Replace missing values operator



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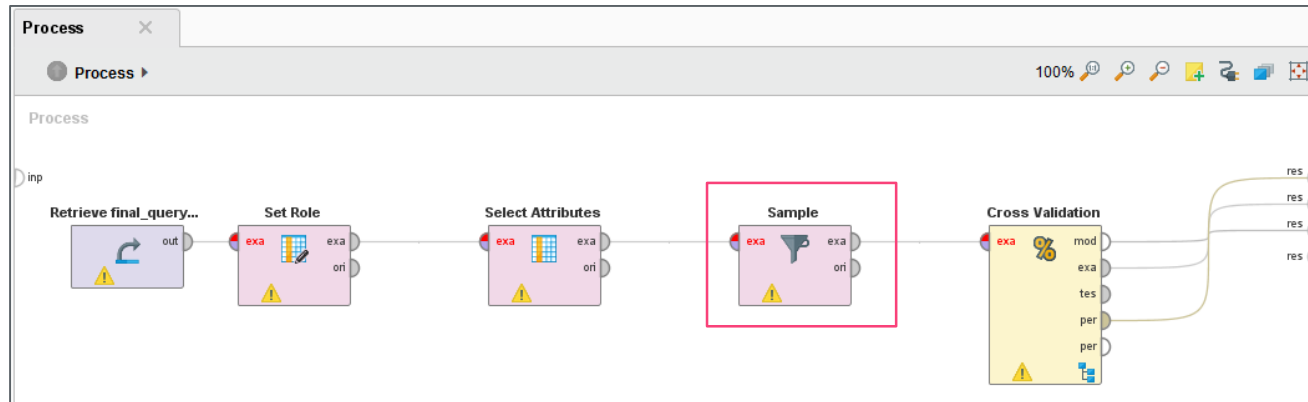
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# Developing a Clinical Prediction Model

## Evaluation and Validation: RapidMiner

- Sample operator: can be used for balancing datasets

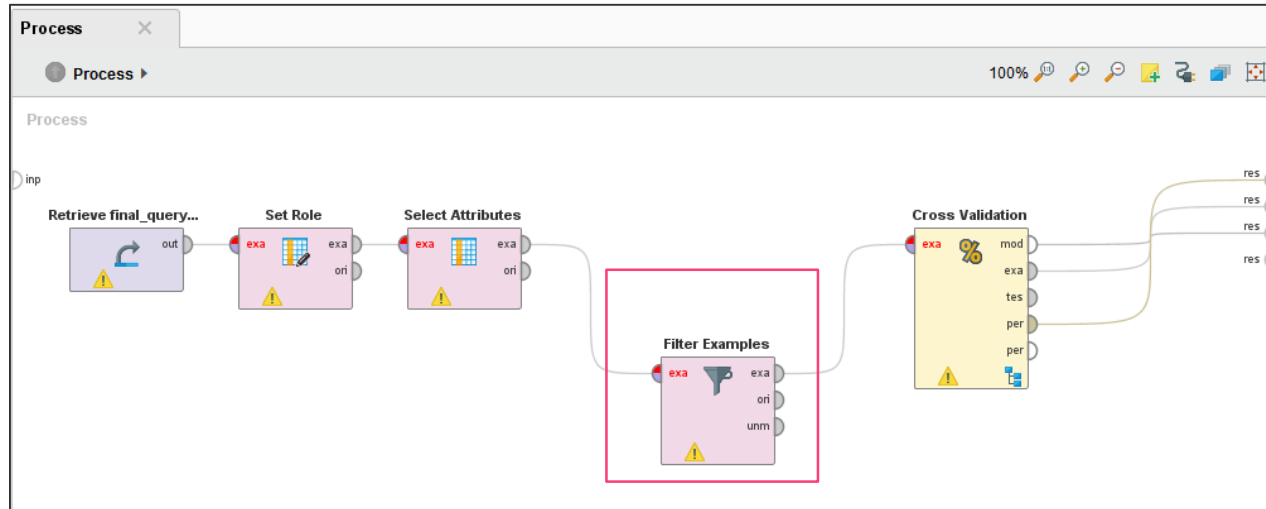


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# Developing a Clinical Prediction Model Evaluation and Validation: RapidMiner

- Filter examples operator: remove all missing data

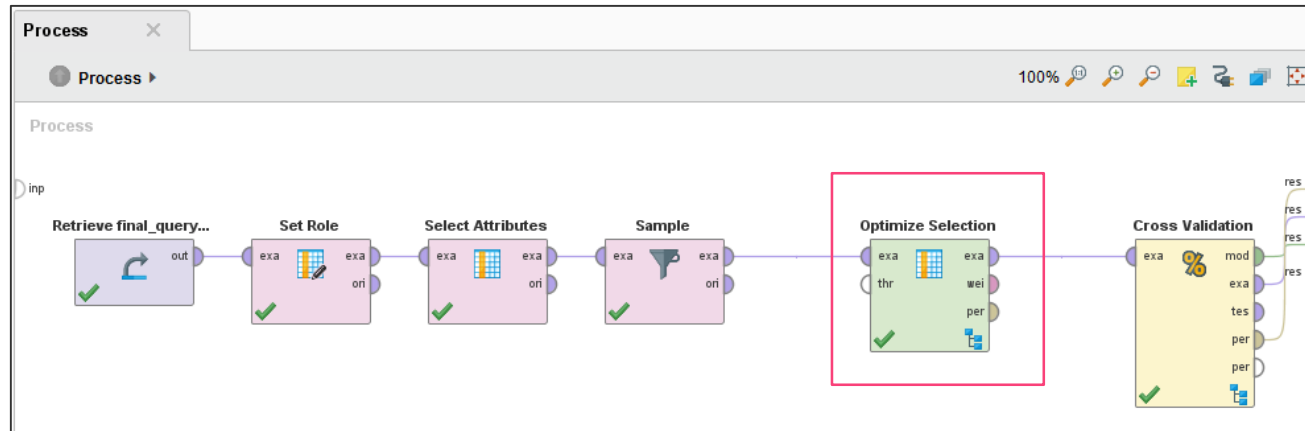


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# Developing a Clinical Prediction Model Evaluation and Validation: RapidMiner

- Optimize selection operator

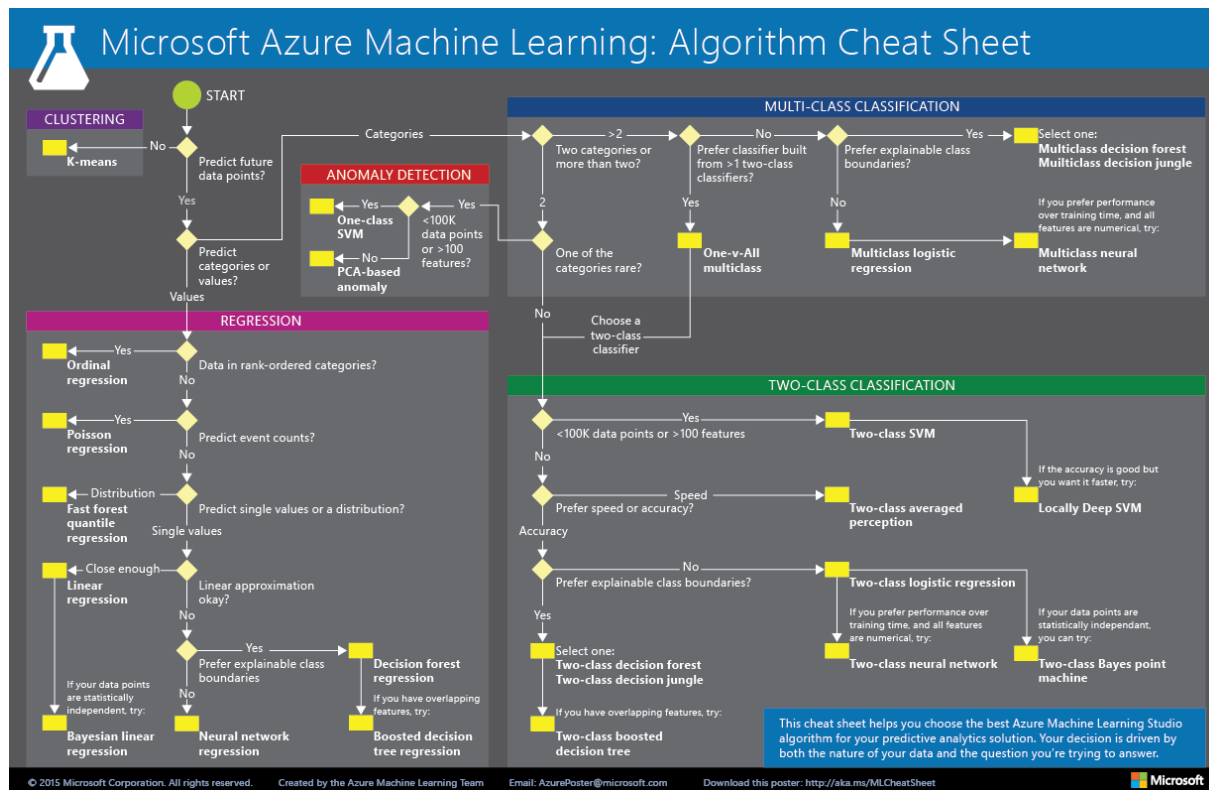


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# Developing a Clinical Prediction Model

## How to choose an algorithm?



<http://oliviaklose.azurewebsites.net/machine-learning-11-algorithms-explained/>

# Exercise II

## Developing a Clinical Prediction Model for Nephrology

- What you will need
  - Download and install RapidMiner
  - Obtain an academic license
  - Install JDBC HANA driver for RapidMiner
  - Connect to HANA and perform experiments
- Proceed according to the step-by-step process
  - Develop the model
  - Answer the questions on openHPI



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# What to Take Home?

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- Developing a clinical prediction model step-by-step
- Impact of missing data on performance of the models
- Impact of class balancing on model performance
- Important concepts: precision, recall, ROC curve
- Evaluating and validating models

# What's Coming Next?

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- Use Case Oncology

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