

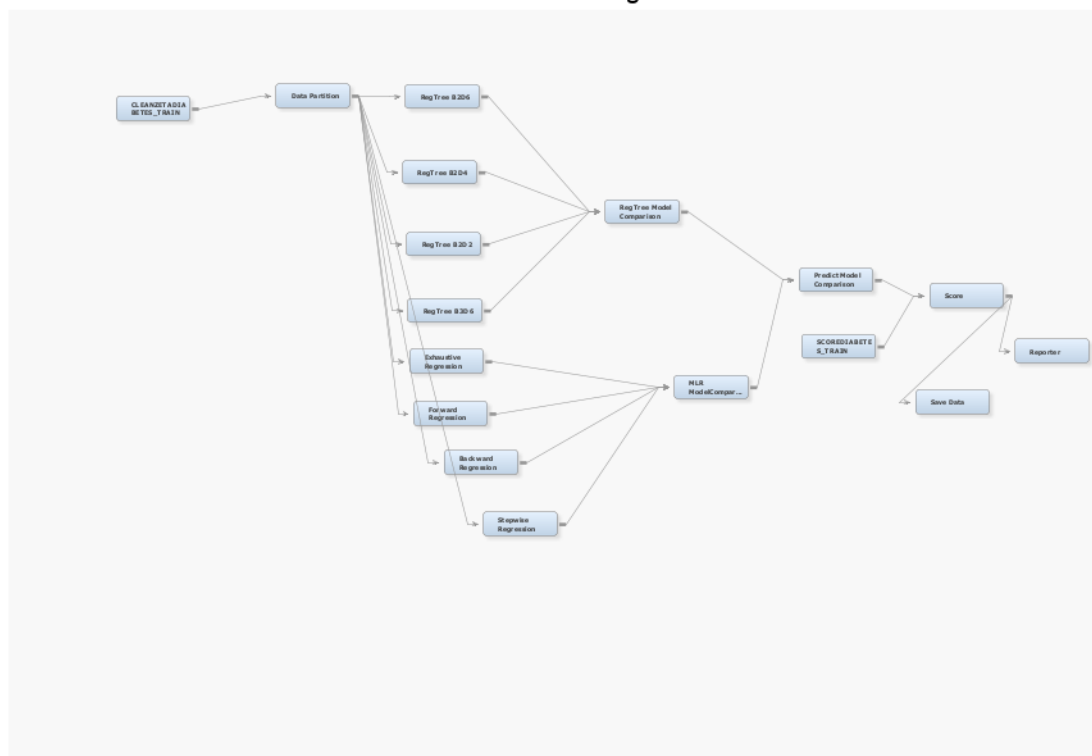
SAS Enterprise Miner Report

User = shre2
 Date = 01:50:07 March 04
 Project = SV.PredictiveAnalysis
 Diagram = SV.Prediction

Start Node = Report
 Node label = Reporter
 Nodes = PATH
 Showall = N

Format = PDF
 Style = LISTING

SAS Enterprise Miner Report Process Flow Diagram



SAS Enterprise Miner Report

Node=CLEANZETADIABETES_TRAIN
Summary

Node id = Ids3
Node label = CLEANZETADIABETES_TRAIN
Meta path = Ids3
Notes =

Node=CLEANZETADIABETES_TRAIN
Properties

| Property | Value | Default | Property | Value | Default |
|------------------------------|----------------------------------|---------|----------------------|------------------------|---------|
| Component | DataSource | | DsCreatedBy | shre2 | |
| ApplyIntervalLevelLowerLimit | Y | | DsId | cleanzetadiabetestrain | |
| ApplyMaxClassLevels | Y | | DsModifiedBy | shre2 | |
| ApplyMaxPercentMissing | Y | | DsModifyDate | 2025129977.8 | |
| CMeta | WORK.M12BVFD5 | | DsSampleName | | |
| ComputeStatistics | N | | DsSampleSize | | |
| DBPassThrough | Y | | DsSampleSizeType | | |
| Data | ZETADATA.CLEANZETADIABETES_TRAIN | | DsScope | LOCAL | |
| DataSelection | DATASOURCE | | IdentifyEmptyColumns | Y | |
| DataSource | cleanzetadiabetestrain | | IntervalLowerLimit | 20 | |
| DataSourceRole | RAW | | Library | ZETADATA | |
| Description | | | MaxClassLevels | 20 | |
| DropMapVariables | Y | | MaxPercentMissing | 50 | |
| DsCreateDate | 2025129977.5 | | MetaAdvisor | BASIC | |

| Property | Value | Default |
|-------------------|-------------------------|---------|
| NBytes | 37094400 | . |
| NCols | 22 | . |
| NObs | 200572 | . |
| NewTable | | |
| NewVariableRole | REJECT | |
| OutputType | VIEW | |
| Role | RAW | TRAIN |
| Sample | D | |
| SampleSizeObs | 10000 | |
| SampleSizePercent | 20 | |
| SampleSizeType | PERCENT | |
| Scope | LOCAL | |
| Segment | | |
| Table | CLEANZETADIABETES_TRAIN | |

Node=CLEANZETADIABETES_TRAIN
Data Attributes

| Attribute | Value | Attribute | Value | Attribute | Value |
|------------|-------------------------|----------------|--------------------|--------------|----------|
| Data Name | CLEANZETADIABETES_TRAIN | Date Created | 04Mar2024:00:00:32 | Data Size | 37094400 |
| Data Type | DATA | Date Modified | 04Mar2024:00:00:32 | Role | RAW |
| Data Label | | Number Rows | 200572 | Segment | |
| Engine | BASE | Number Columns | 22 | Data Library | ZETADATA |

Node=CLEANZETADIABETES_TRAIN
Variables List

| Name | Label | Role | Level | Type | Length | Format | Creator |
|----------------------|----------------------|--------|----------|------|--------|--------|---------|
| Age | Age | INPUT | INTERVAL | N | 8 | BEST. | |
| AnyHealthcare | AnyHealthcare | INPUT | BINARY | N | 8 | BEST. | |
| BMI | BMI | INPUT | INTERVAL | N | 8 | BEST. | |
| CholCheck | CholCheck | INPUT | BINARY | N | 8 | BEST. | |
| Diabetes_Prediction | | TARGET | NOMINAL | C | 11 | | |
| DiffWalk | DiffWalk | INPUT | BINARY | N | 8 | BEST. | |
| Education | Education | INPUT | ORDINAL | N | 8 | BEST. | |
| Fruits | Fruits | INPUT | BINARY | N | 8 | BEST. | |
| GenHlth | GenHlth | INPUT | ORDINAL | N | 8 | BEST. | |
| HeartDiseaseorAttack | HeartDiseaseorAttack | INPUT | BINARY | N | 8 | BEST. | |
| HighBP | HighBP | INPUT | BINARY | N | 8 | BEST. | |
| HighChol | HighChol | INPUT | BINARY | N | 8 | BEST. | |
| HvyAlcoholConsump | HvyAlcoholConsump | INPUT | BINARY | N | 8 | BEST. | |
| Income | Income | INPUT | ORDINAL | N | 8 | BEST. | |
| MentHlth | MentHlth | INPUT | ORDINAL | N | 8 | BEST. | |
| NoDocbcCost | NoDocbcCost | INPUT | BINARY | N | 8 | BEST. | |
| PhysActivity | PhysActivity | INPUT | BINARY | N | 8 | BEST. | |
| PhysHlth | PhysHlth | INPUT | ORDINAL | N | 8 | BEST. | |
| Sex | Sex | INPUT | BINARY | N | 8 | BEST. | |
| Smoker | Smoker | INPUT | BINARY | N | 8 | BEST. | |
| Stroke | Stroke | INPUT | BINARY | N | 8 | BEST. | |
| Veggies | Veggies | INPUT | BINARY | N | 8 | BEST. | |

SAS Enterprise Miner Report

Node=Data Partition Summary

Node id = Part
Node label = Data Partition
Meta path = Ids3 => Part
Notes =

Node=Data Partition Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|----------------------|-----------|---------|------------|---------|---------|-------------|-------|---------|
| Component | Partition | | Method | DEFAULT | | TestPct | 30 | |
| ClassDistribution | Y | | OutputType | DATA | | TrainPct | 40 | |
| IntervalDistribution | Y | | RandomSeed | 12345 | | ValidatePct | 30 | |

Node=Data Partition Variable Summary

| Role | Level | Frequency Count | Name |
|--------|----------|--------------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |

SAS Enterprise Miner Report

Node=Exhaustive Regression
Summary

Node id = Reg
Node label = Exhaustive Regression
Meta path = Ids3 => Part => Reg
Notes =

Node=Exhaustive Regression
Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|------------------|--------------|--------------|-----------------------|-----------|---------|--------------------|---------|---------|
| Component | Regression | | Force | 0 | | PolynomialDegree | 2 | |
| AbsConvValue | -1.34078E154 | -7.237006E75 | GConvTimes | 1 | | PrintDesignMatrix | N | |
| AbsFTime | 1 | | GConvValue | 1E-6 | | Rule | NONE | |
| AbsFValue | 0 | | Hierarchy | CLASS | | SASSPDS | N | |
| AbsGTime | 1 | | InputCoding | DEVIATION | | SelectionCriterion | DEFAULT | |
| AbsGValue | 0.00001 | | Interactions | | | SelectionDefault | Y | |
| AbsXTime | 1 | | LinkFunction | LOGIT | | Sequential | N | |
| AbsXValue | 1E-8 | | MainEffect | Y | | Simple | N | |
| CIParam | N | | MaxCPUTime | 1 HOUR | | SIEntry | 0.05 | |
| ConvDefaults | Y | | MaxFunctionCalls | . | | SIStay | 0.05 | |
| CorB | N | | MaxIterations | . | | Start | 0 | |
| CovB | N | | MaxStep | . | | StepOutput | N | |
| Covout | N | | MinResourceUse | N | | Stop | 0 | |
| Details | N | | ModelDefaults | Y | | SuppressIntercept | N | |
| Error | LOGISTIC | | ModelSelection | NONE | | SuppressOutput | N | |
| ExcludedVariable | REJECT | | OptimizationTechnique | DEFAULT | | Terms | N | |
| FConvTimes | 1 | | Performance | N | | TwoFactor | N | |
| FConvValue | 0 | | Polynomial | N | | | | |

Node=Exhaustive Regression
Variable Summary

| Role | Level | Frequency Count | Name |
|--------|----------|-----------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |

Node=Exhaustive Regression
Model Fit Statistics

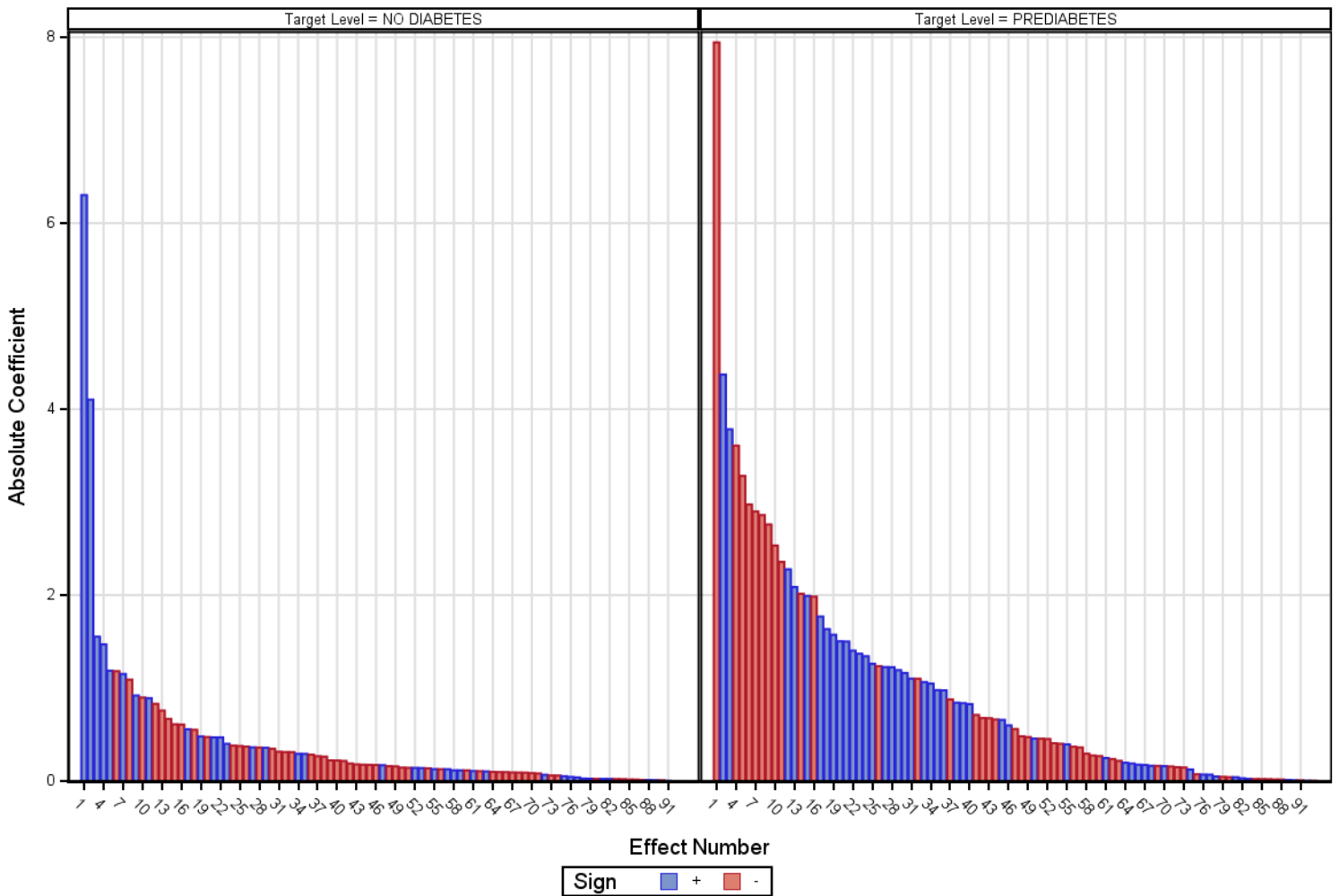
Target=Diabetes_Prediction Target Label=' '

| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|------|
| Akaike's Information Criterion | 63558.08 | . | . |
| Average Squared Error | 0.07 | 0.07 | 0.07 |
| Average Error Function | 0.26 | 0.26 | 0.26 |
| Degrees of Freedom for Error | 160268.00 | . | . |
| Model Degrees of Freedom | 186.00 | . | . |

Target=Diabetes_Prediction Target Label=' '

| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|-----------|
| Total Degrees of Freedom | 160454.00 | . | . |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Error Function | 63186.08 | 47745.74 | 47539.06 |
| Final Prediction Error | 0.07 | . | . |
| Maximum Absolute Error | 1.00 | 1.00 | 1.00 |
| Mean Square Error | 0.07 | 0.07 | 0.07 |
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Number of Estimate Weights | 186.00 | . | . |
| Root Average Sum of Squares | 0.27 | 0.27 | 0.27 |
| Root Final Prediction Error | 0.27 | . | . |
| Root Mean Squared Error | 0.27 | 0.27 | 0.27 |
| Schwarz's Bayesian Criterion | 65415.43 | . | . |
| Sum of Squared Errors | 17714.06 | 13322.68 | 13303.93 |
| Sum of Case Weights Times Freq | 240681.00 | 180507.00 | 180528.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |

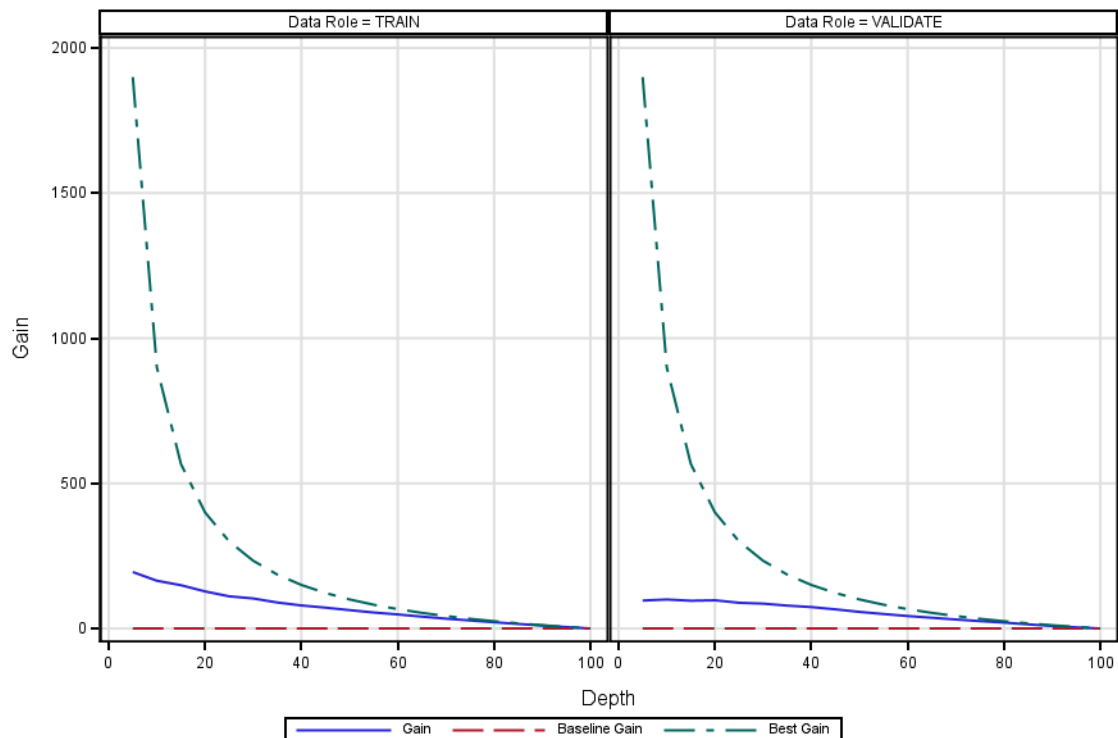
SAS Enterprise Miner Report
Node=Exhaustive Regression
Regression Model Effects



SAS Enterprise Miner Report

Node=Exhaustive Regression

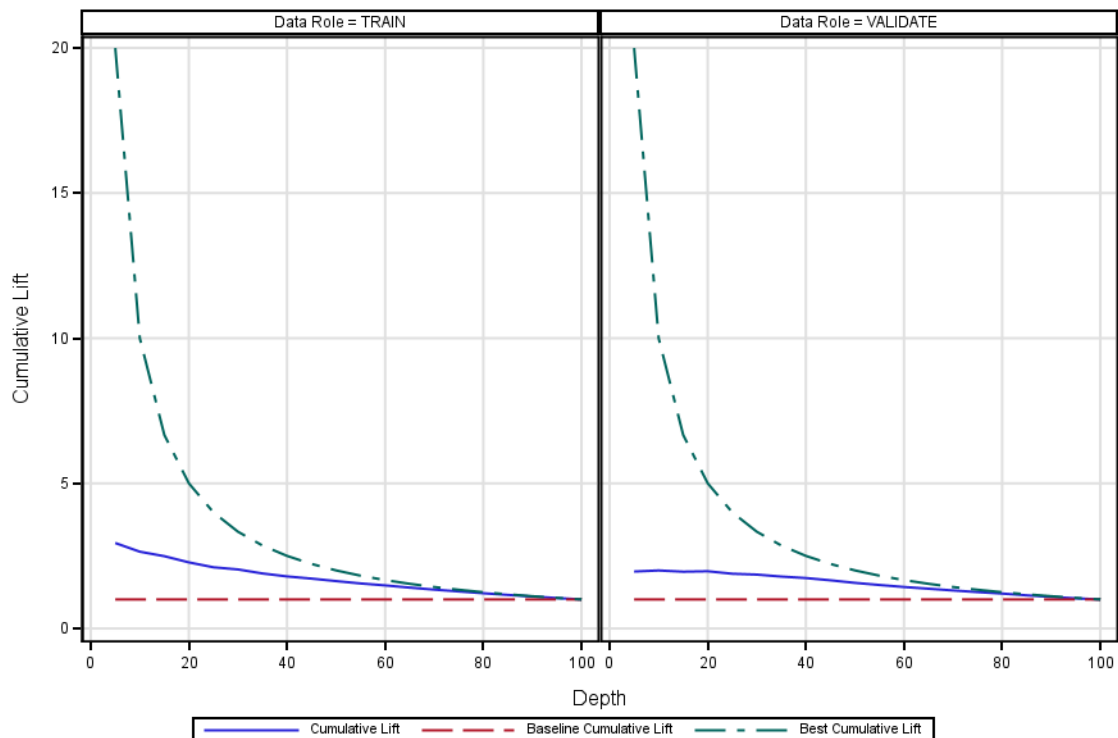
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=Exhaustive Regression

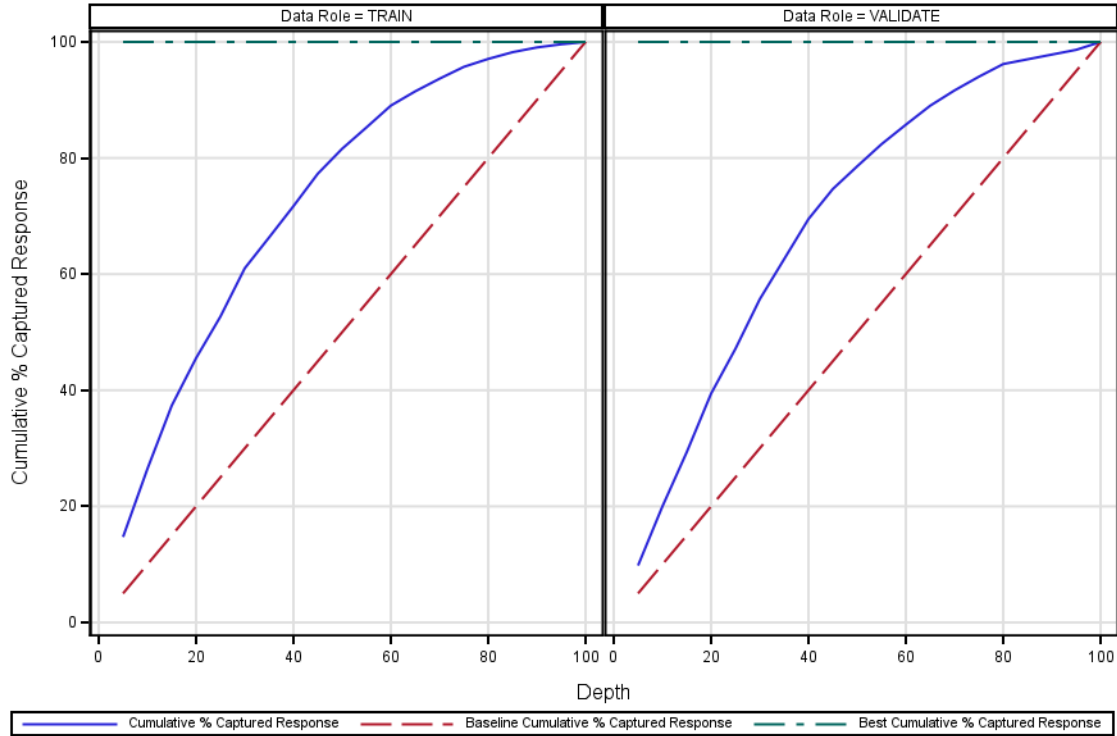
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

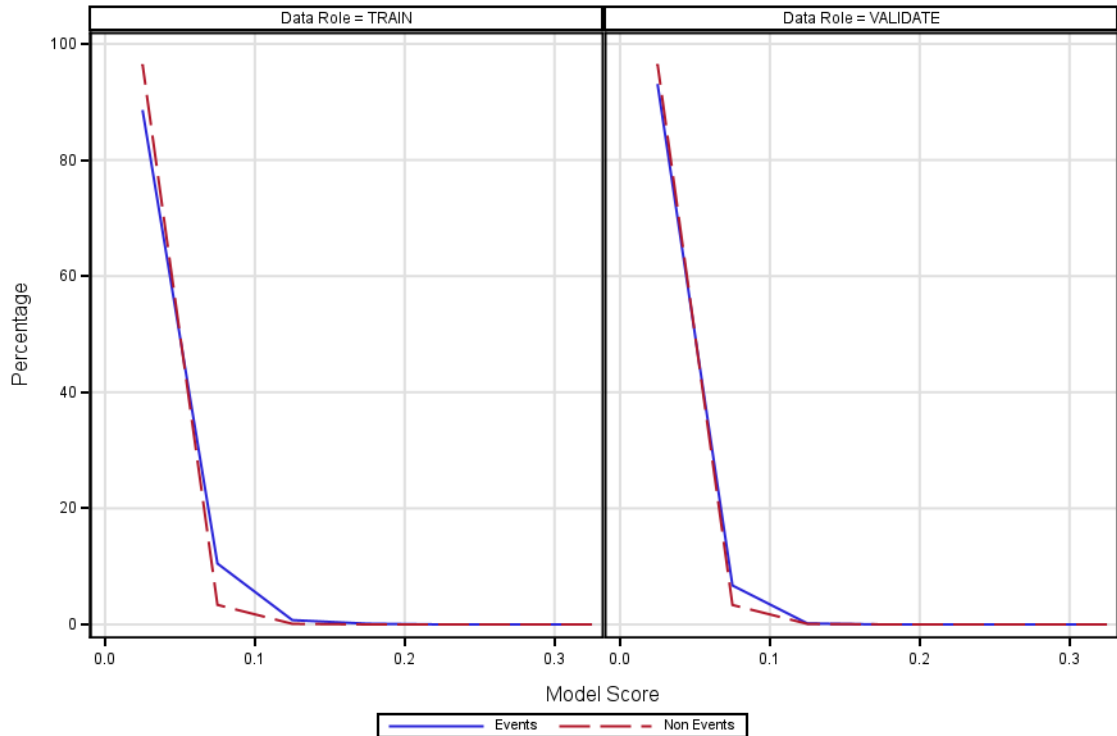
Node=Exhaustive Regression

Model Assessment Scores where TARGET='Diabetes_Prediction'

**SAS Enterprise Miner Report**

Node=Exhaustive Regression

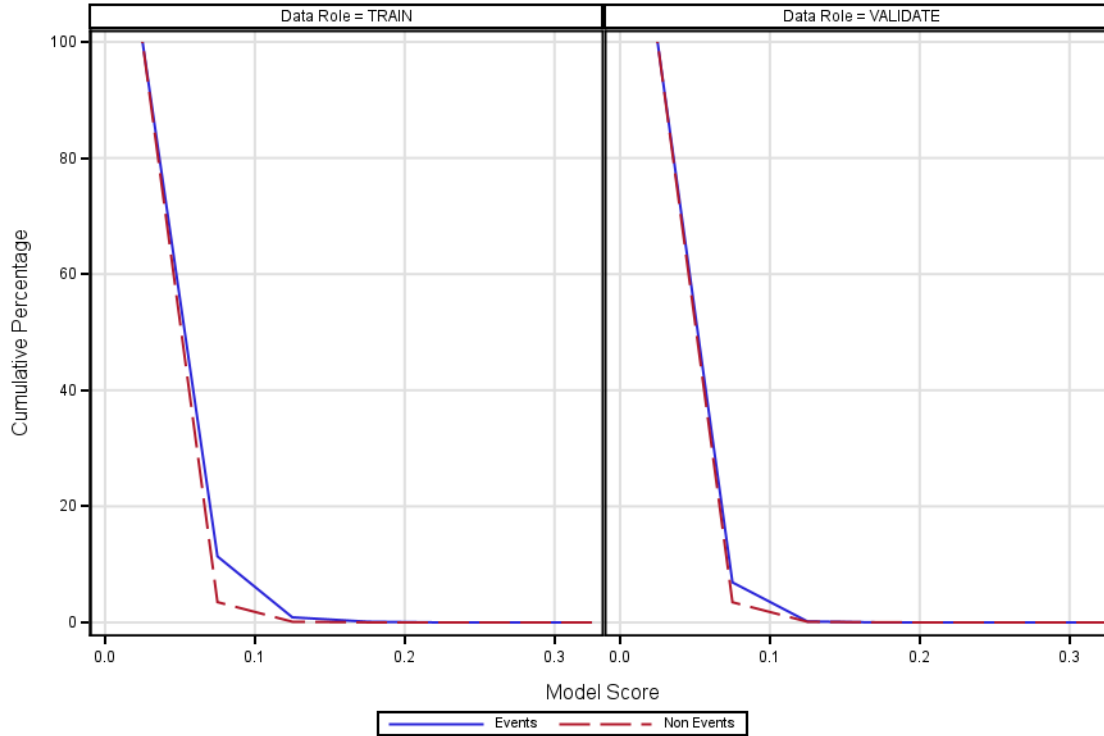
Score Distributions where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=Exhaustive Regression

Score Distributions where TARGET='Diabetes_Prediction'



Node=Exhaustive Regression Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.30-0.35 | 0 | 0.0000 | 0.0025 | 0.000 | 0.003 |
| 0.25-0.30 | 0 | 0.0000 | 0.0013 | 0.000 | 0.004 |
| 0.20-0.25 | 0 | 0.0000 | 0.0038 | 0.000 | 0.008 |
| 0.15-0.20 | 2 | 0.1363 | 0.0102 | 0.136 | 0.018 |
| 0.10-0.15 | 11 | 0.7498 | 0.1117 | 0.886 | 0.130 |
| 0.05-0.10 | 154 | 10.4976 | 3.3761 | 11.384 | 3.506 |
| 0.00-0.05 | 1300 | 88.6162 | 96.4944 | 100.000 | 100.000 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.30-0.35 | 0 | 0.0000 | 0.0034 | 0.000 | 0.003 |
| 0.25-0.30 | 0 | 0.0000 | 0.0017 | 0.000 | 0.005 |
| 0.20-0.25 | 0 | 0.0000 | 0.0051 | 0.000 | 0.010 |
| 0.15-0.20 | 0 | 0.0000 | 0.0135 | 0.000 | 0.024 |
| 0.10-0.15 | 2 | 0.1817 | 0.1016 | 0.182 | 0.125 |
| 0.05-0.10 | 74 | 6.7212 | 3.3571 | 6.903 | 3.482 |
| 0.00-0.05 | 1025 | 93.0972 | 96.5176 | 100.000 | 100.000 |

SAS Enterprise Miner Report

Node=Forward Regression Summary

Node id = Reg2
Node label = Forward Regression
Meta path = Ids3 => Part => Reg2
Notes =

Node=Forward Regression Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|------------------|--------------|--------------|-----------------------|-----------|---------|--------------------|---------|---------|
| Component | Regression | | Force | 0 | | PolynomialDegree | 2 | |
| AbsConvValue | -1.34078E154 | -7.237006E75 | GConvTimes | 1 | | PrintDesignMatrix | N | |
| AbsFTime | 1 | | GConvValue | 1E-6 | | Rule | NONE | |
| AbsFValue | 0 | | Hierarchy | CLASS | | SASSPDS | N | |
| AbsGTime | 1 | | InputCoding | DEVIATION | | SelectionCriterion | DEFAULT | |
| AbsGValue | 0.00001 | | Interactions | | | SelectionDefault | Y | |
| AbsXTime | 1 | | LinkFunction | LOGIT | | Sequential | N | |
| AbsXValue | 1E-8 | | MainEffect | Y | | Simple | N | |
| CIParam | N | | MaxCPUTime | 1 HOUR | | SIEntry | 0.05 | |
| ConvDefaults | Y | | MaxFunctionCalls | . | | SIStay | 0.05 | |
| CorB | N | | MaxIterations | . | | Start | 0 | |
| CovB | N | | MaxStep | . | | StepOutput | N | |
| Covout | N | | MinResourceUse | N | | Stop | 0 | |
| Details | N | | ModelDefaults | Y | | SuppressIntercept | N | |
| Error | LOGISTIC | | ModelSelection | FORWARD | NONE | SuppressOutput | N | |
| ExcludedVariable | REJECT | | OptimizationTechnique | DEFAULT | | Terms | N | |
| FConvTimes | 1 | | Performance | N | | TwoFactor | N | |
| FConvValue | 0 | | Polynomial | N | | | | |

Node=Forward Regression Variable Summary

| Role | Level | Frequency Count | Name |
|--------|----------|-----------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |

Node=Forward Regression Model Fit Statistics

Target=Diabetes_Prediction Target Label=' '

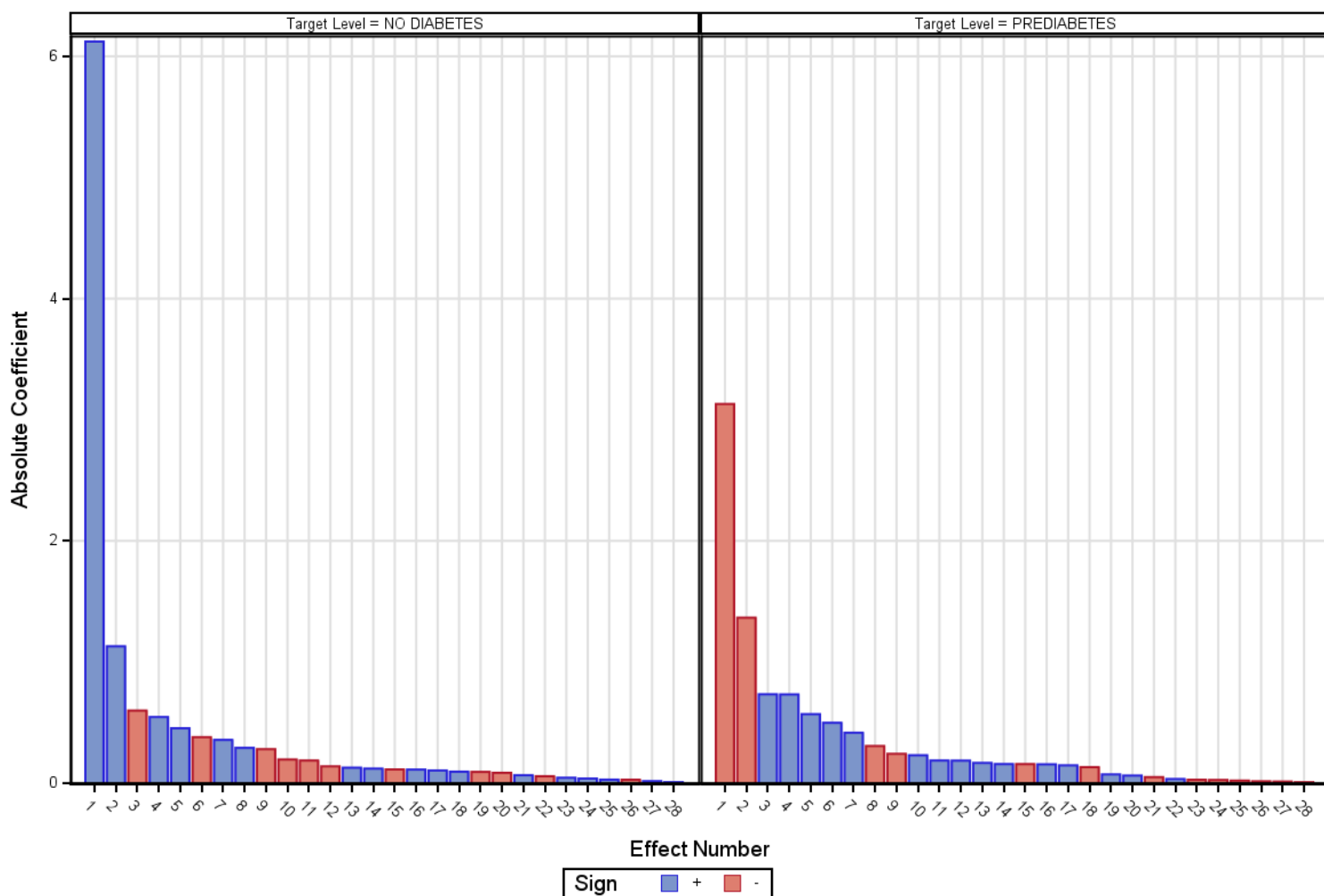
| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|------|
| Akaike's Information Criterion | 63457.86 | . | . |
| Average Squared Error | 0.07 | 0.07 | 0.07 |
| Average Error Function | 0.26 | 0.26 | 0.26 |
| Degrees of Freedom for Error | 160398.00 | . | . |
| Model Degrees of Freedom | 56.00 | . | . |

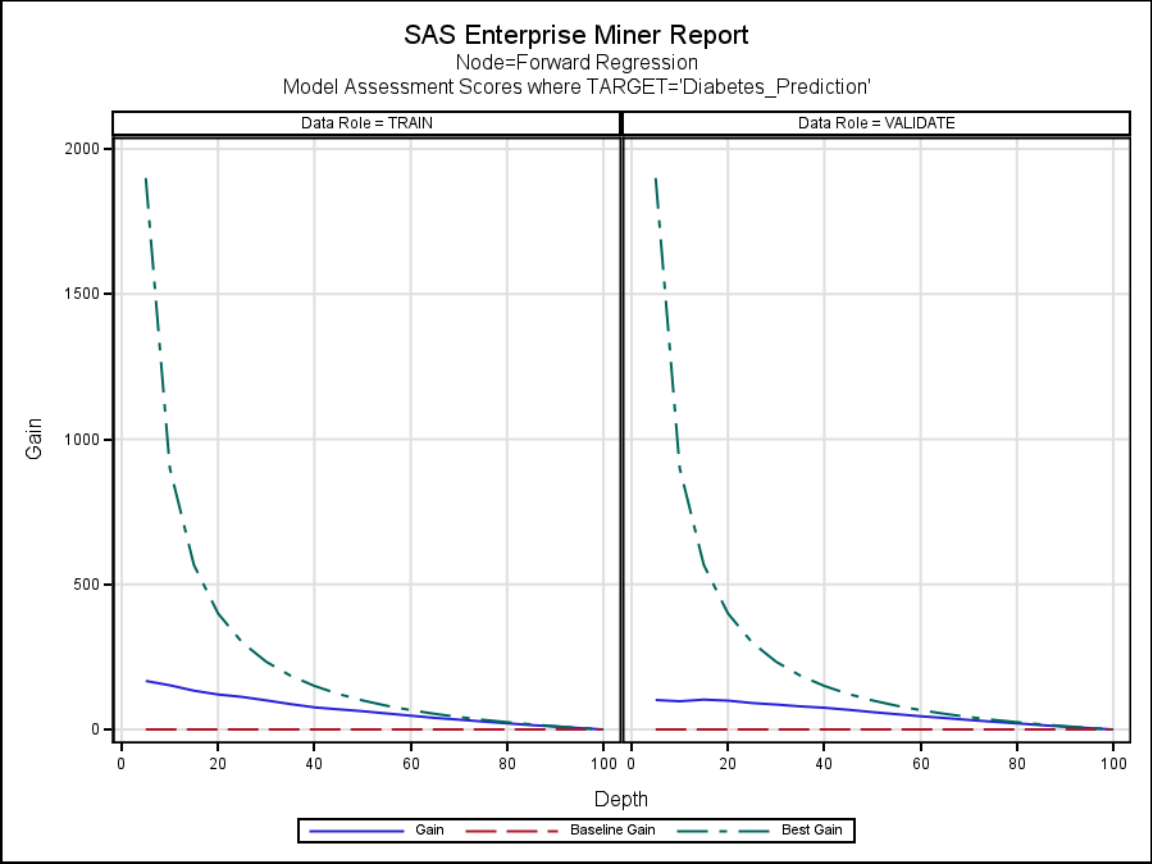
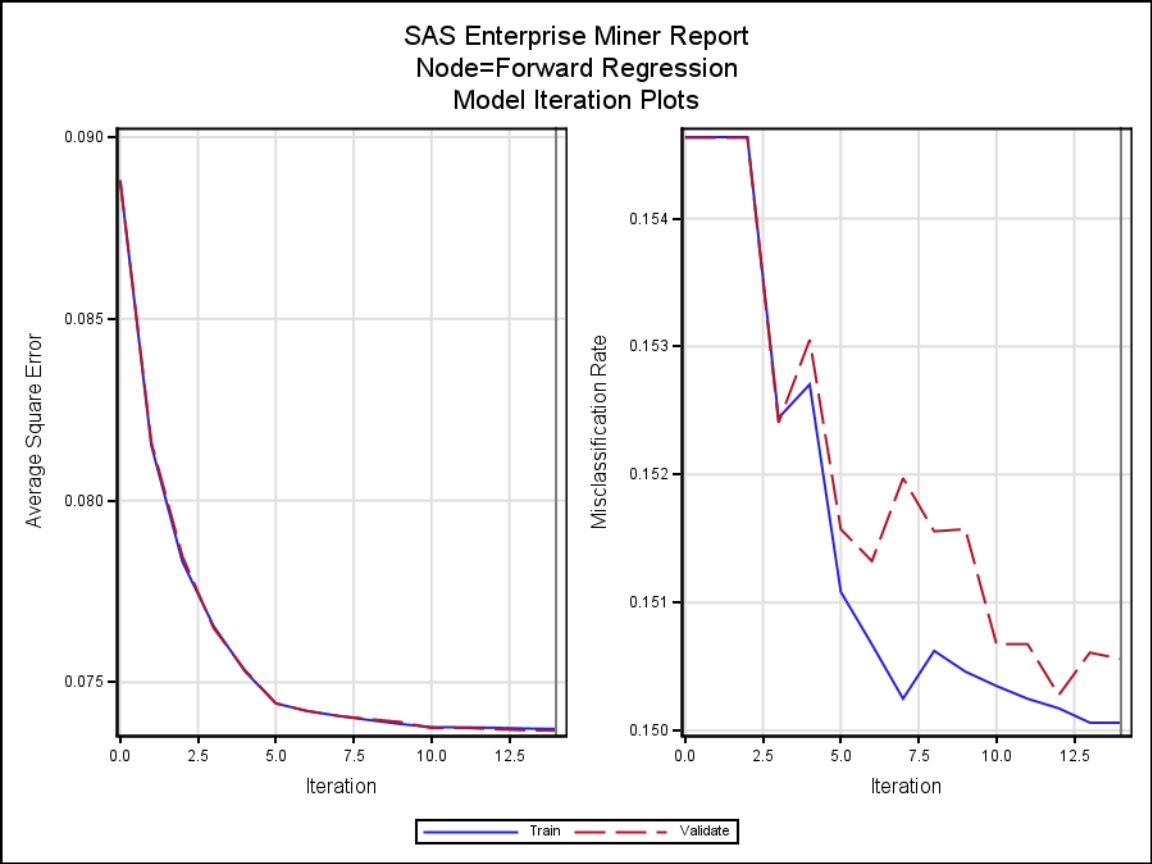
Target=Diabetes_Prediction Target Label=' '

| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|-----------|
| Total Degrees of Freedom | 160454.00 | . | . |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Error Function | 63345.86 | 47578.87 | 47439.71 |
| Final Prediction Error | 0.07 | . | . |
| Maximum Absolute Error | 1.00 | 1.00 | 1.00 |
| Mean Square Error | 0.07 | 0.07 | 0.07 |
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Number of Estimate Weights | 56.00 | . | . |
| Root Average Sum of Squares | 0.27 | 0.27 | 0.27 |
| Root Final Prediction Error | 0.27 | . | . |
| Root Mean Squared Error | 0.27 | 0.27 | 0.27 |
| Schwarz's Bayesian Criterion | 64017.07 | . | . |
| Sum of Squared Errors | 17742.41 | 13298.36 | 13290.51 |
| Sum of Case Weights Times Freq | 240681.00 | 180507.00 | 180528.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |

SAS Enterprise Miner Report

Node=Forward Regression
Regression Model Effects

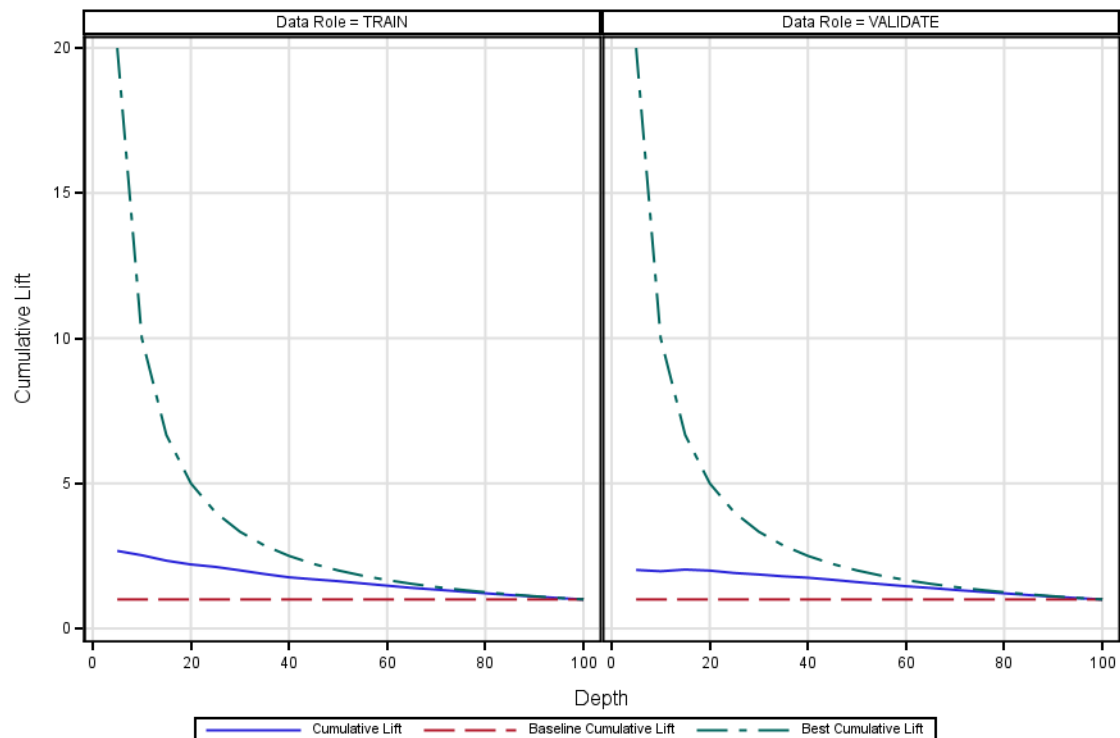




SAS Enterprise Miner Report

Node=Forward Regression

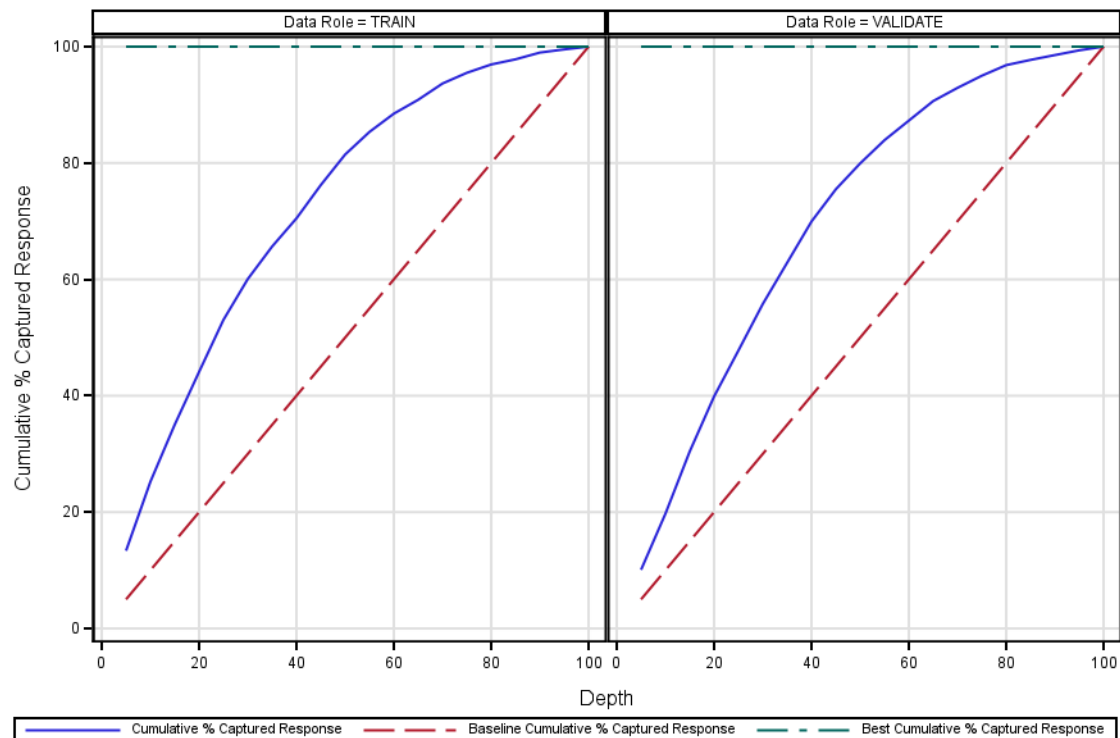
Model Assessment Scores where TARGET='Diabetes_Prediction'

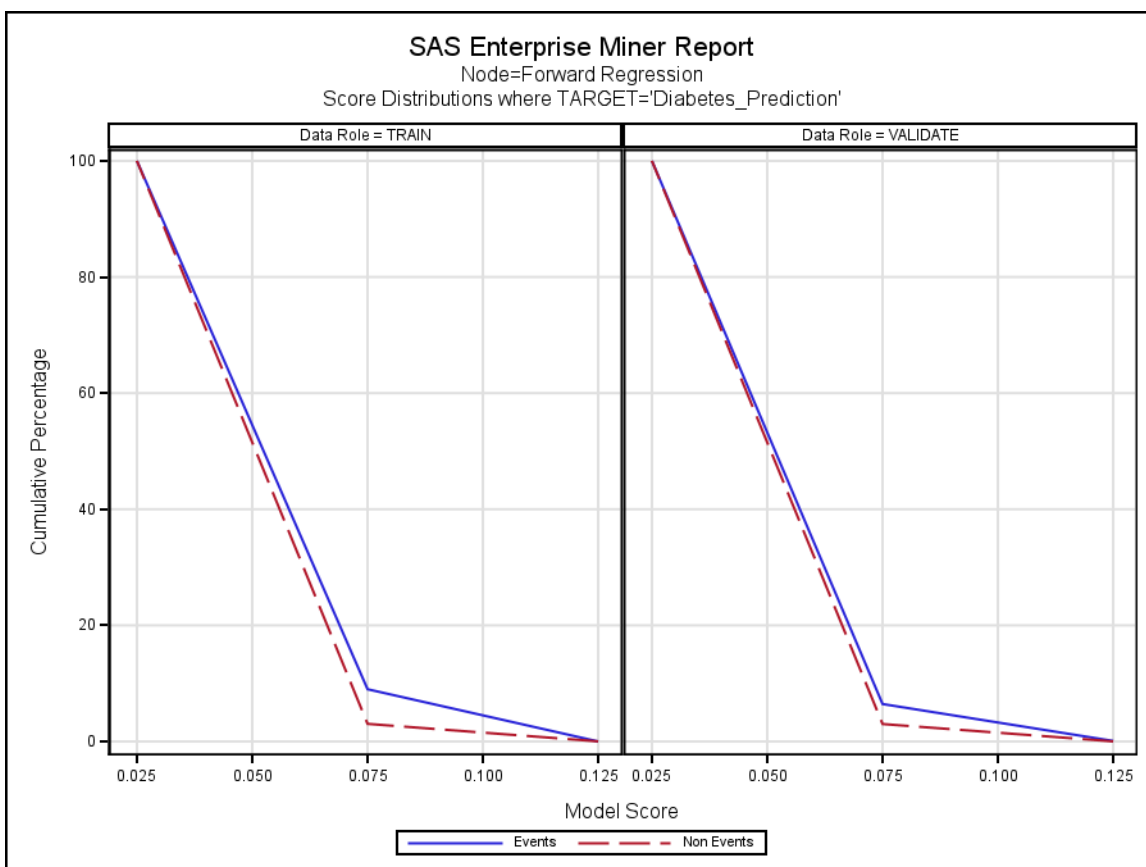
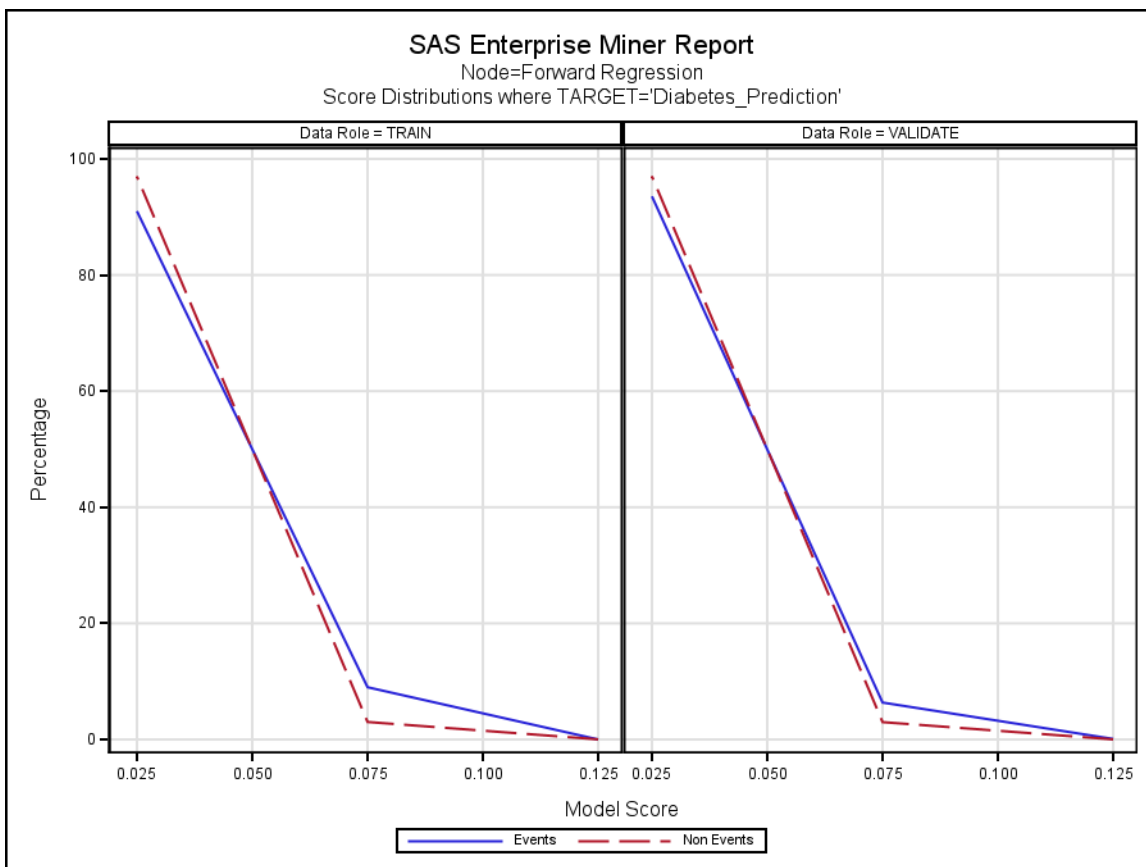


SAS Enterprise Miner Report

Node=Forward Regression

Model Assessment Scores where TARGET='Diabetes_Prediction'





Node=Forward Regression
Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.10-0.15 | 0 | 0.0000 | 0.0203 | 0.000 | 0.020 |
| 0.05-0.10 | 132 | 8.9980 | 3.0142 | 8.998 | 3.035 |
| 0.00-0.05 | 1335 | 91.0020 | 96.9655 | 100.000 | 100.000 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.10-0.15 | 1 | 0.0908 | 0.0169 | 0.091 | 0.017 |
| 0.05-0.10 | 70 | 6.3579 | 2.9881 | 6.449 | 3.005 |
| 0.00-0.05 | 1030 | 93.5513 | 96.9950 | 100.000 | 100.000 |

SAS Enterprise Miner Report

Node=Backward Regression Summary

Node id = Reg3
Node label = Backward Regression
Meta path = Ids3 => Part => Reg3
Notes =

Node=Backward Regression Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|------------------|--------------|--------------|-----------------------|-----------|---------|--------------------|---------|---------|
| Component | Regression | | Force | 0 | | PolynomialDegree | 2 | |
| AbsConvValue | -1.34078E154 | -7.237006E75 | GConvTimes | 1 | | PrintDesignMatrix | N | |
| AbsFTime | 1 | | GConvValue | 1E-6 | | Rule | NONE | |
| AbsFValue | 0 | | Hierarchy | CLASS | | SASSPDS | N | |
| AbsGTime | 1 | | InputCoding | DEVIATION | | SelectionCriterion | DEFAULT | |
| AbsGValue | 0.00001 | | Interactions | | | SelectionDefault | Y | |
| AbsXTime | 1 | | LinkFunction | LOGIT | | Sequential | N | |
| AbsXValue | 1E-8 | | MainEffect | Y | | Simple | N | |
| CIParam | N | | MaxCPUTime | 1 HOUR | | SIEntry | 0.05 | |
| ConvDefaults | Y | | MaxFunctionCalls | . | | SIStay | 0.05 | |
| CorB | N | | MaxIterations | . | | Start | 0 | |
| CovB | N | | MaxStep | . | | StepOutput | N | |
| Covout | N | | MinResourceUse | N | | Stop | 0 | |
| Details | N | | ModelDefaults | Y | | SuppressIntercept | N | |
| Error | LOGISTIC | | ModelSelection | BACKWARD | NONE | SuppressOutput | N | |
| ExcludedVariable | REJECT | | OptimizationTechnique | DEFAULT | | Terms | N | |
| FConvTimes | 1 | | Performance | N | | TwoFactor | N | |
| FConvValue | 0 | | Polynomial | N | | | | |

Node=Backward Regression Variable Summary

| Role | Level | Frequency Count | Name |
|--------|----------|-----------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |

Node=Backward Regression Model Fit Statistics

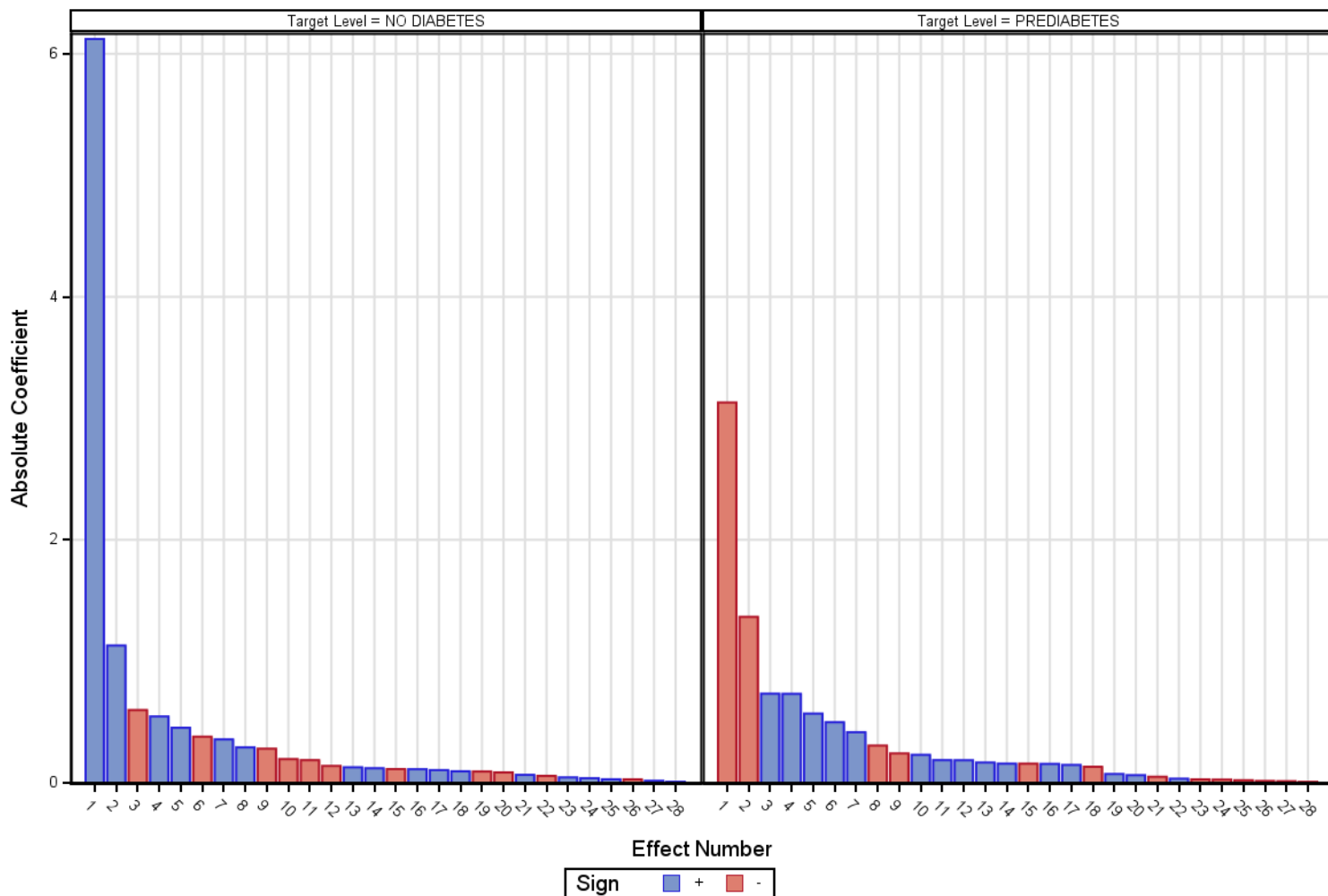
Target=Diabetes_Prediction Target Label=' '

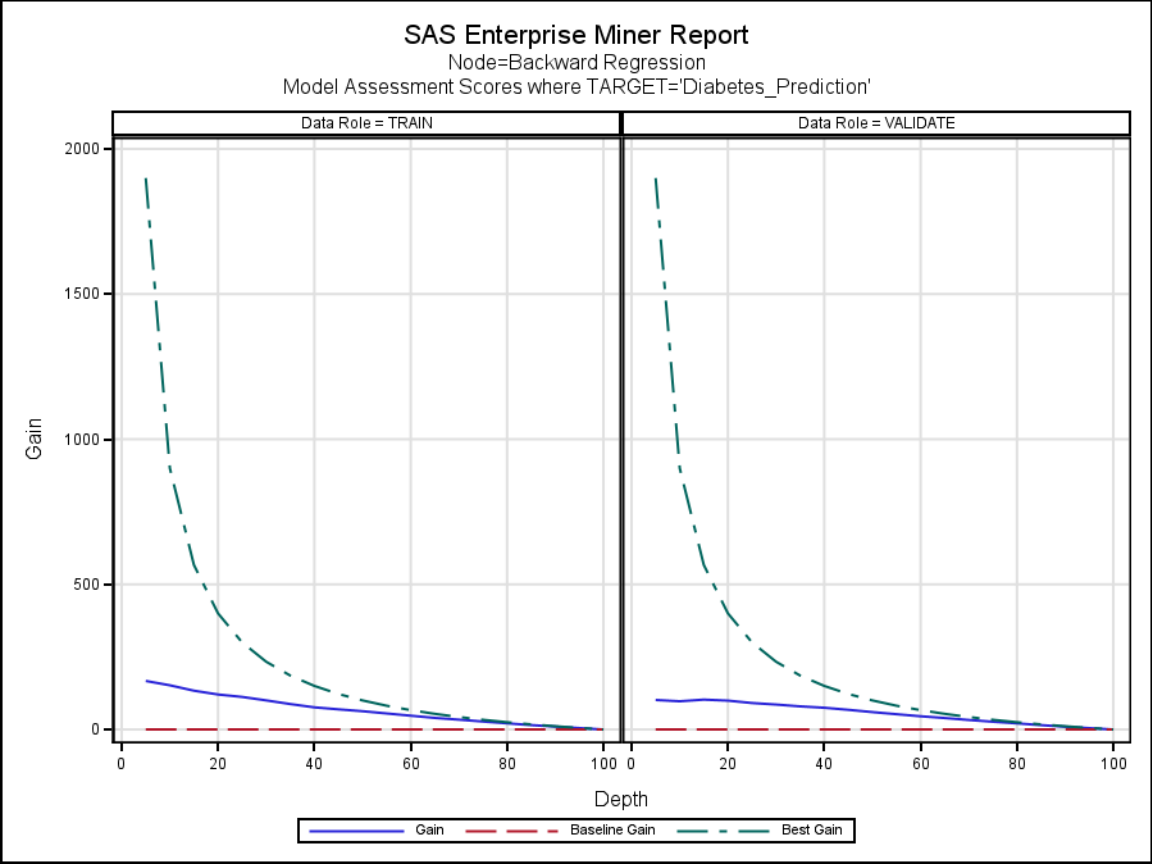
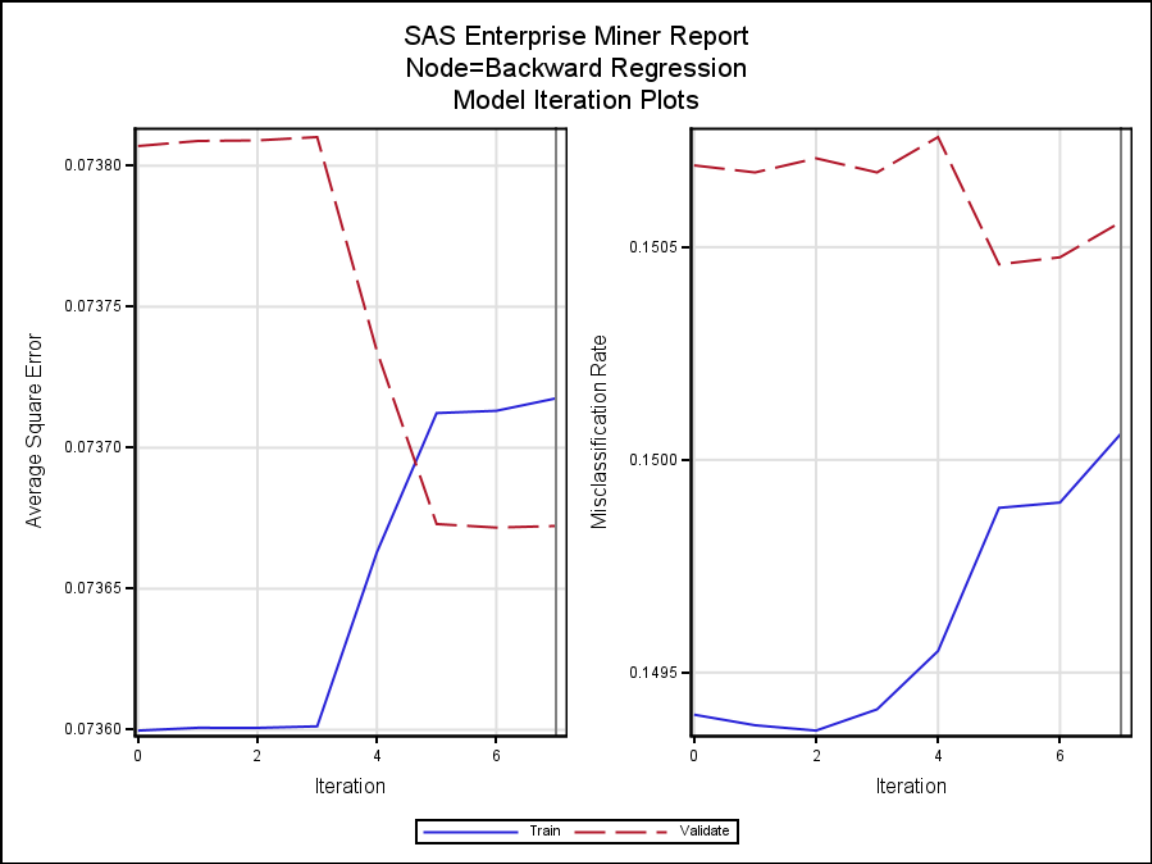
| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|------|
| Akaike's Information Criterion | 63457.86 | . | . |
| Average Squared Error | 0.07 | 0.07 | 0.07 |
| Average Error Function | 0.26 | 0.26 | 0.26 |
| Degrees of Freedom for Error | 160398.00 | . | . |
| Model Degrees of Freedom | 56.00 | . | . |

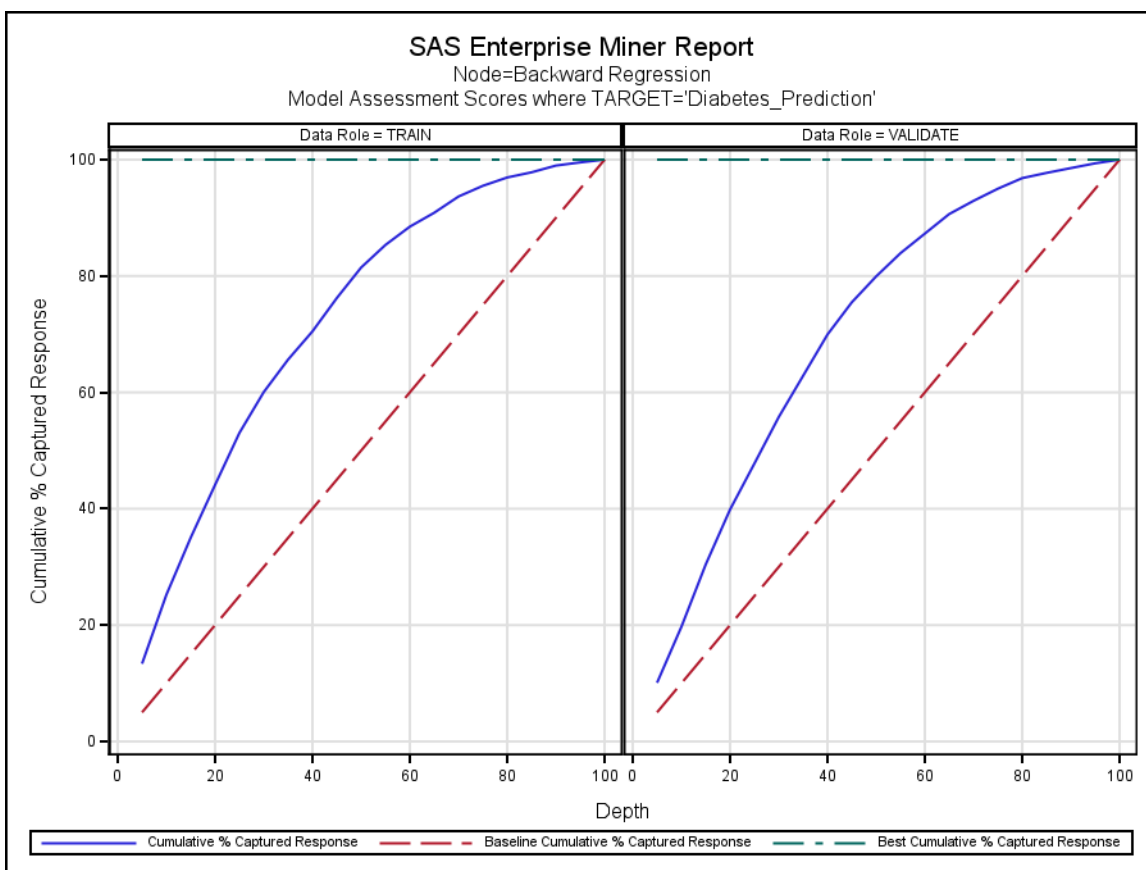
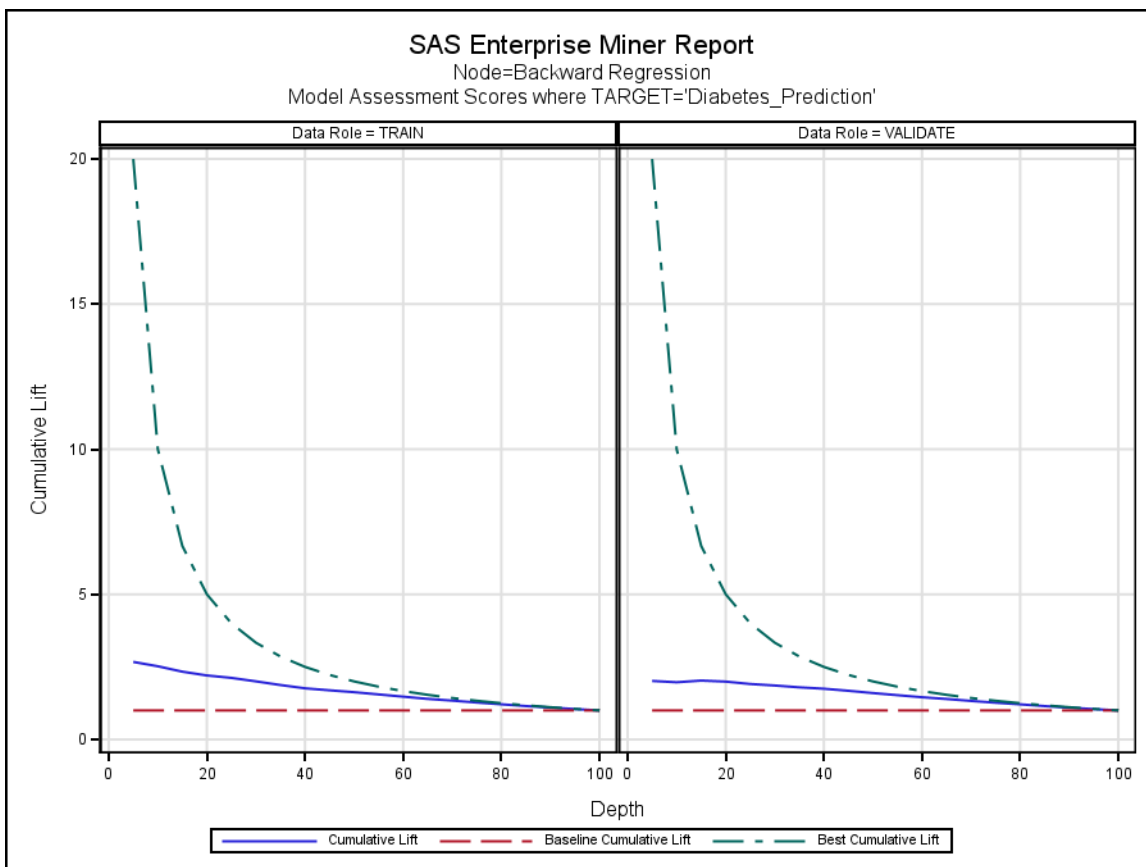
Target=Diabetes_Prediction Target Label=' '

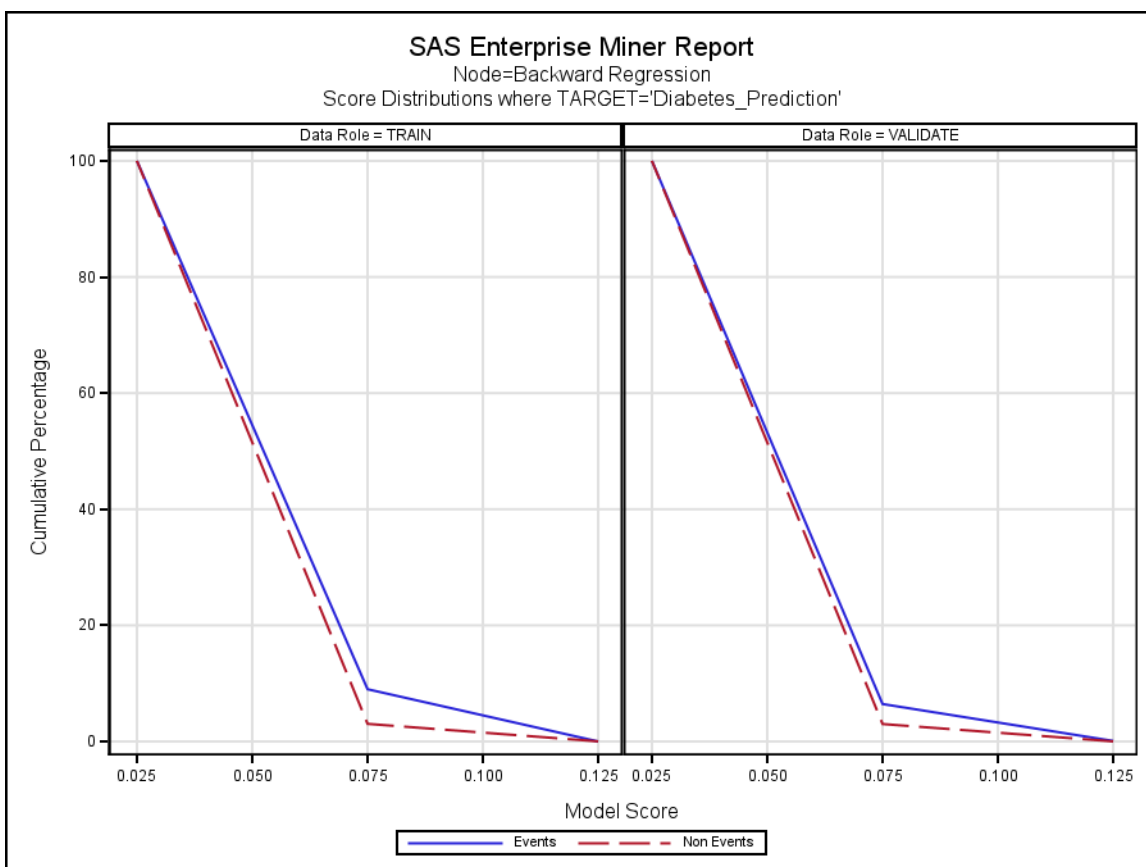
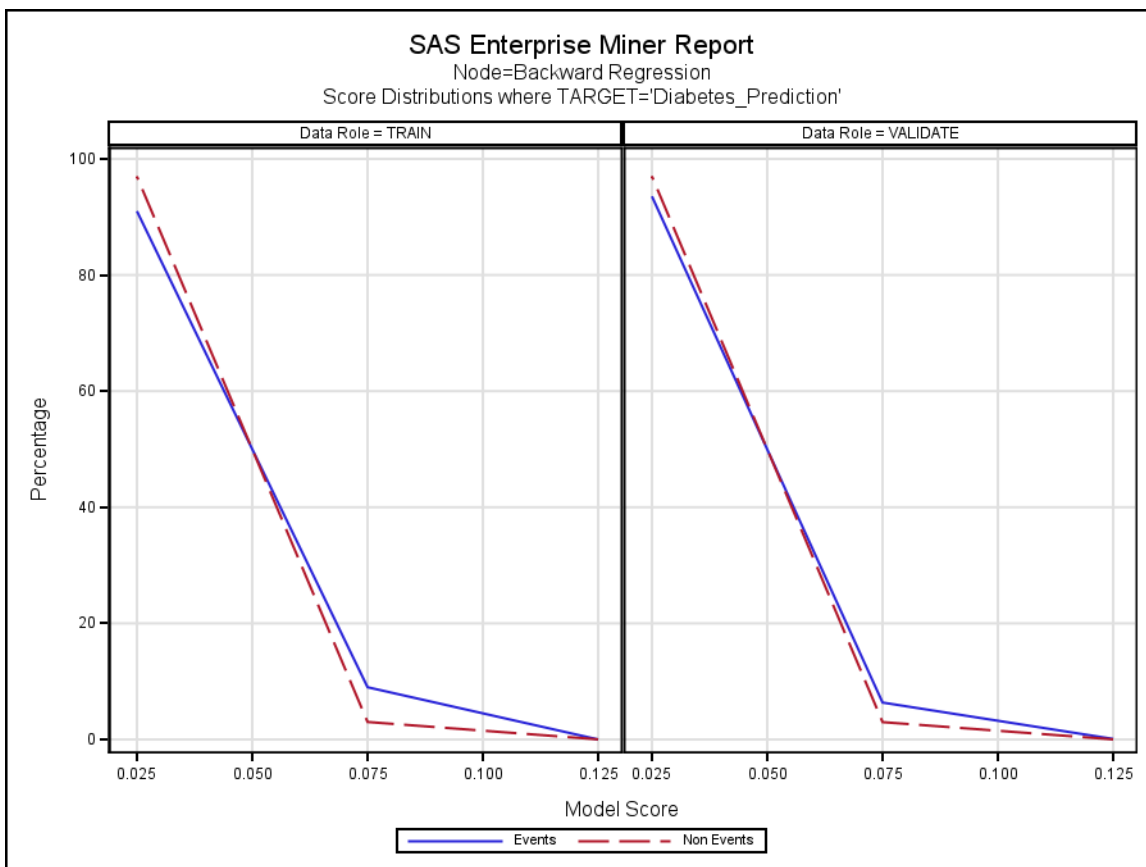
| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|-----------|
| Total Degrees of Freedom | 160454.00 | . | . |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Error Function | 63345.86 | 47578.87 | 47439.71 |
| Final Prediction Error | 0.07 | . | . |
| Maximum Absolute Error | 1.00 | 1.00 | 1.00 |
| Mean Square Error | 0.07 | 0.07 | 0.07 |
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Number of Estimate Weights | 56.00 | . | . |
| Root Average Sum of Squares | 0.27 | 0.27 | 0.27 |
| Root Final Prediction Error | 0.27 | . | . |
| Root Mean Squared Error | 0.27 | 0.27 | 0.27 |
| Schwarz's Bayesian Criterion | 64017.07 | . | . |
| Sum of Squared Errors | 17742.41 | 13298.36 | 13290.51 |
| Sum of Case Weights Times Freq | 240681.00 | 180507.00 | 180528.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |

SAS Enterprise Miner Report
Node=Backward Regression
Regression Model Effects









Node=Backward Regression
Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.10-0.15 | 0 | 0.0000 | 0.0203 | 0.000 | 0.020 |
| 0.05-0.10 | 132 | 8.9980 | 3.0142 | 8.998 | 3.035 |
| 0.00-0.05 | 1335 | 91.0020 | 96.9655 | 100.000 | 100.000 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.10-0.15 | 1 | 0.0908 | 0.0169 | 0.091 | 0.017 |
| 0.05-0.10 | 70 | 6.3579 | 2.9881 | 6.449 | 3.005 |
| 0.00-0.05 | 1030 | 93.5513 | 96.9950 | 100.000 | 100.000 |

SAS Enterprise Miner Report

Node=Stepwise Regression Summary

Node id = Reg4
Node label = Stepwise Regression
Meta path = Ids3 => Part => Reg4
Notes =

Node=Stepwise Regression Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|------------------|--------------|--------------|-----------------------|-----------|---------|--------------------|---------|---------|
| Component | Regression | | Force | 0 | | PolynomialDegree | 2 | |
| AbsConvValue | -1.34078E154 | -7.237006E75 | GConvTimes | 1 | | PrintDesignMatrix | N | |
| AbsFTime | 1 | | GConvValue | 1E-6 | | Rule | NONE | |
| AbsFValue | 0 | | Hierarchy | CLASS | | SASSPDS | N | |
| AbsGTime | 1 | | InputCoding | DEVIATION | | SelectionCriterion | DEFAULT | |
| AbsGValue | 0.00001 | | Interactions | | | SelectionDefault | Y | |
| AbsXTime | 1 | | LinkFunction | LOGIT | | Sequential | N | |
| AbsXValue | 1E-8 | | MainEffect | Y | | Simple | N | |
| CIParam | N | | MaxCPUTime | 1 HOUR | | SIEntry | 0.05 | |
| ConvDefaults | Y | | MaxFunctionCalls | . | | SIStay | 0.05 | |
| CorB | N | | MaxIterations | . | | Start | 0 | |
| CovB | N | | MaxStep | . | | StepOutput | N | |
| Covout | N | | MinResourceUse | N | | Stop | 0 | |
| Details | N | | ModelDefaults | Y | | SuppressIntercept | N | |
| Error | LOGISTIC | | ModelSelection | STEPWISE | NONE | SuppressOutput | N | |
| ExcludedVariable | REJECT | | OptimizationTechnique | DEFAULT | | Terms | N | |
| FConvTimes | 1 | | Performance | N | | TwoFactor | N | |
| FConvValue | 0 | | Polynomial | N | | | | |

Node=Stepwise Regression Variable Summary

| Role | Level | Frequency Count | Name |
|--------|----------|-----------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |

Node=Stepwise Regression Model Fit Statistics

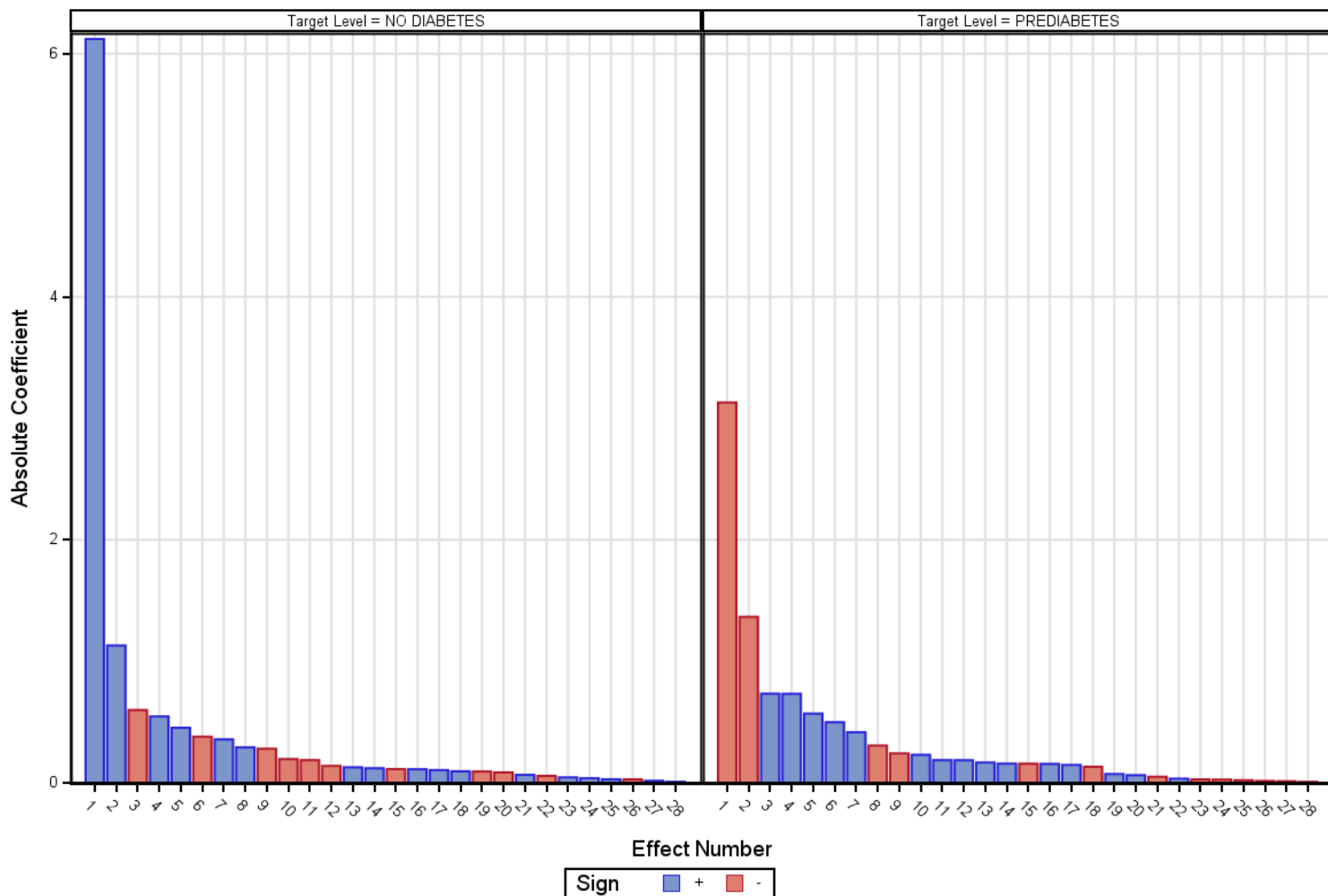
Target=Diabetes_Prediction Target Label=' '

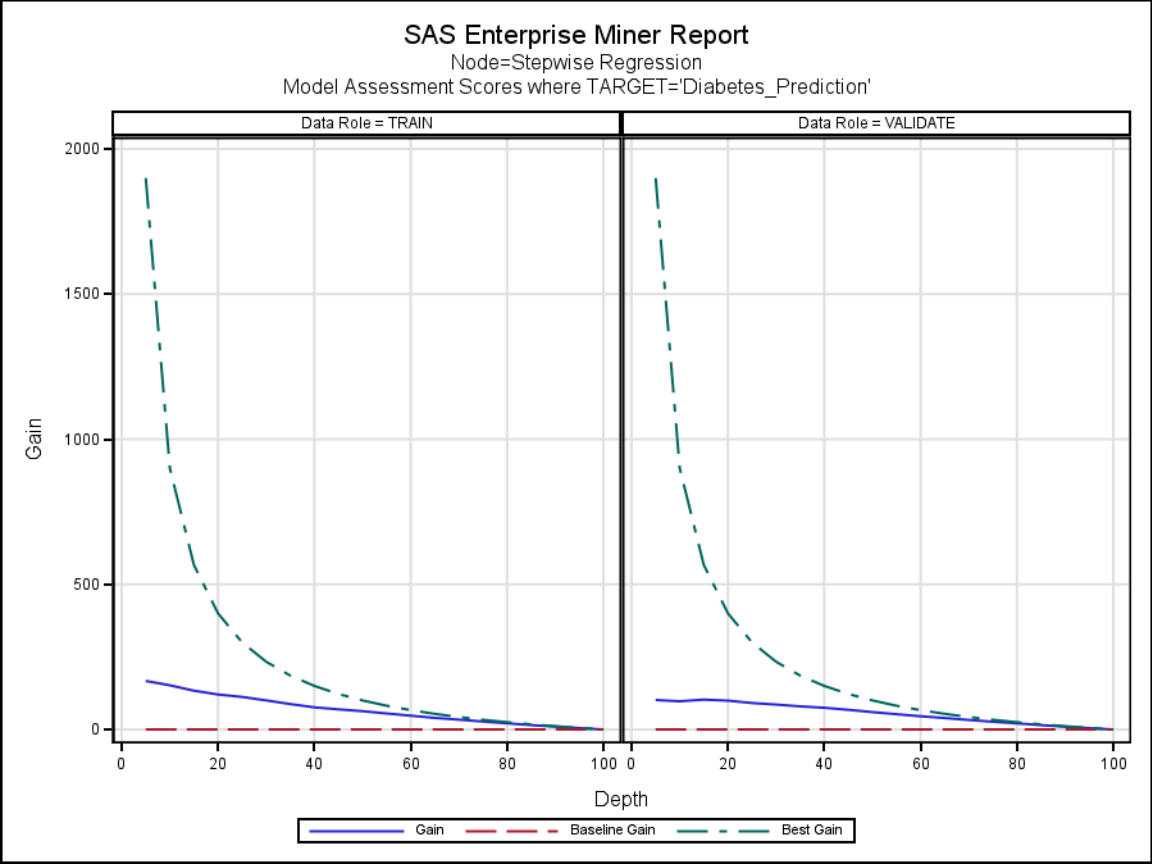
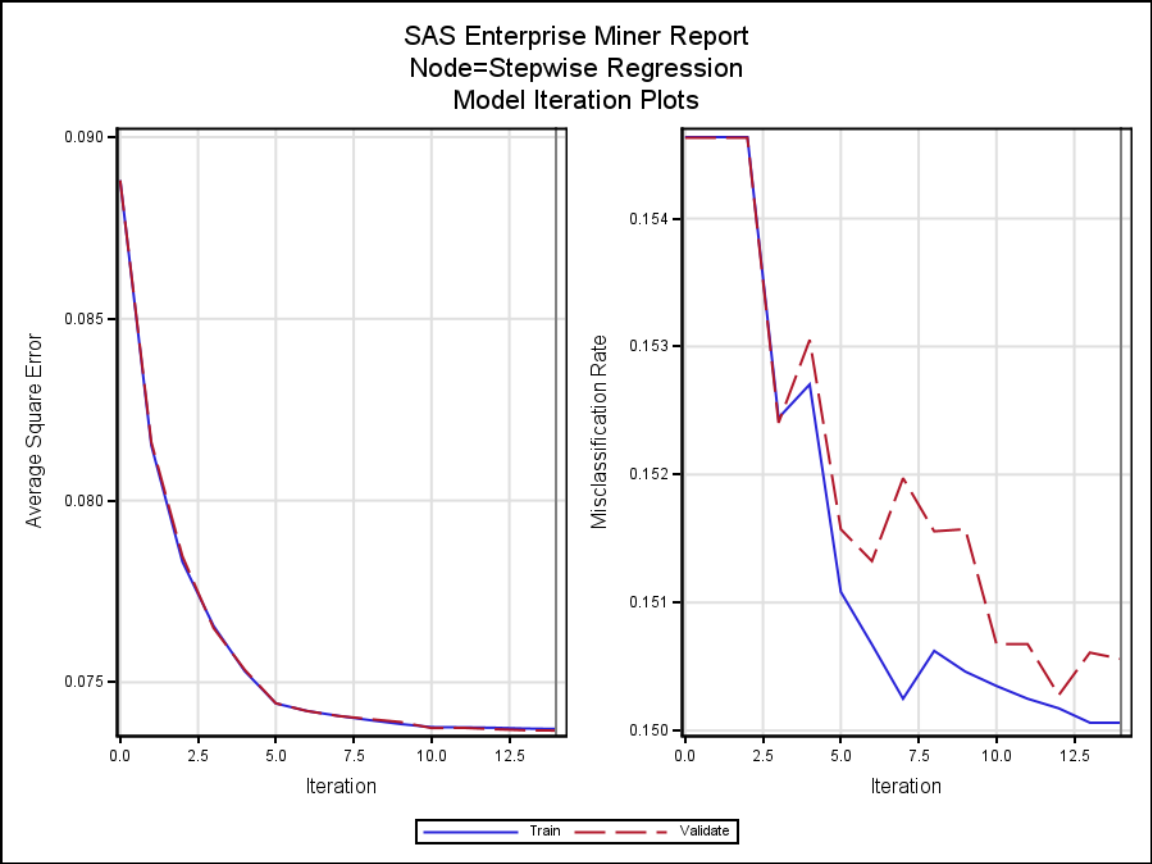
| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|------|
| Akaike's Information Criterion | 63457.86 | . | . |
| Average Squared Error | 0.07 | 0.07 | 0.07 |
| Average Error Function | 0.26 | 0.26 | 0.26 |
| Degrees of Freedom for Error | 160398.00 | . | . |
| Model Degrees of Freedom | 56.00 | . | . |

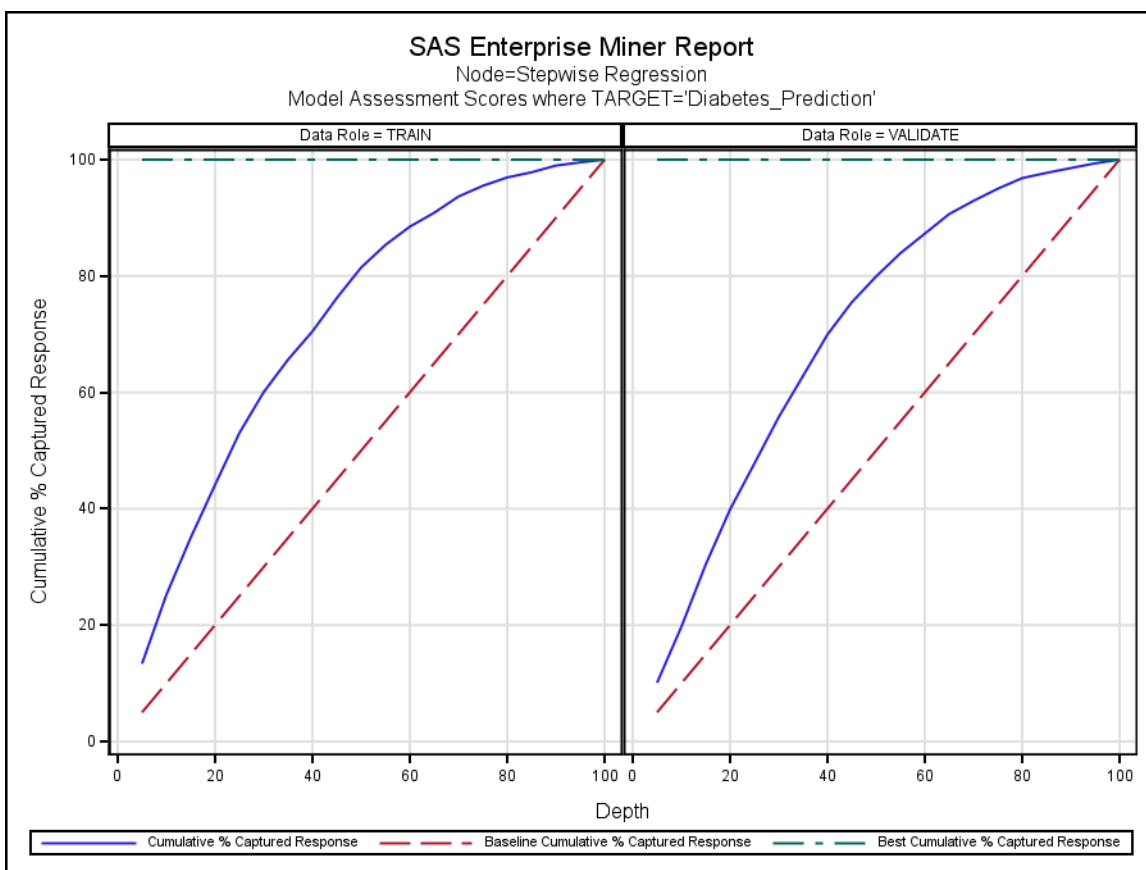
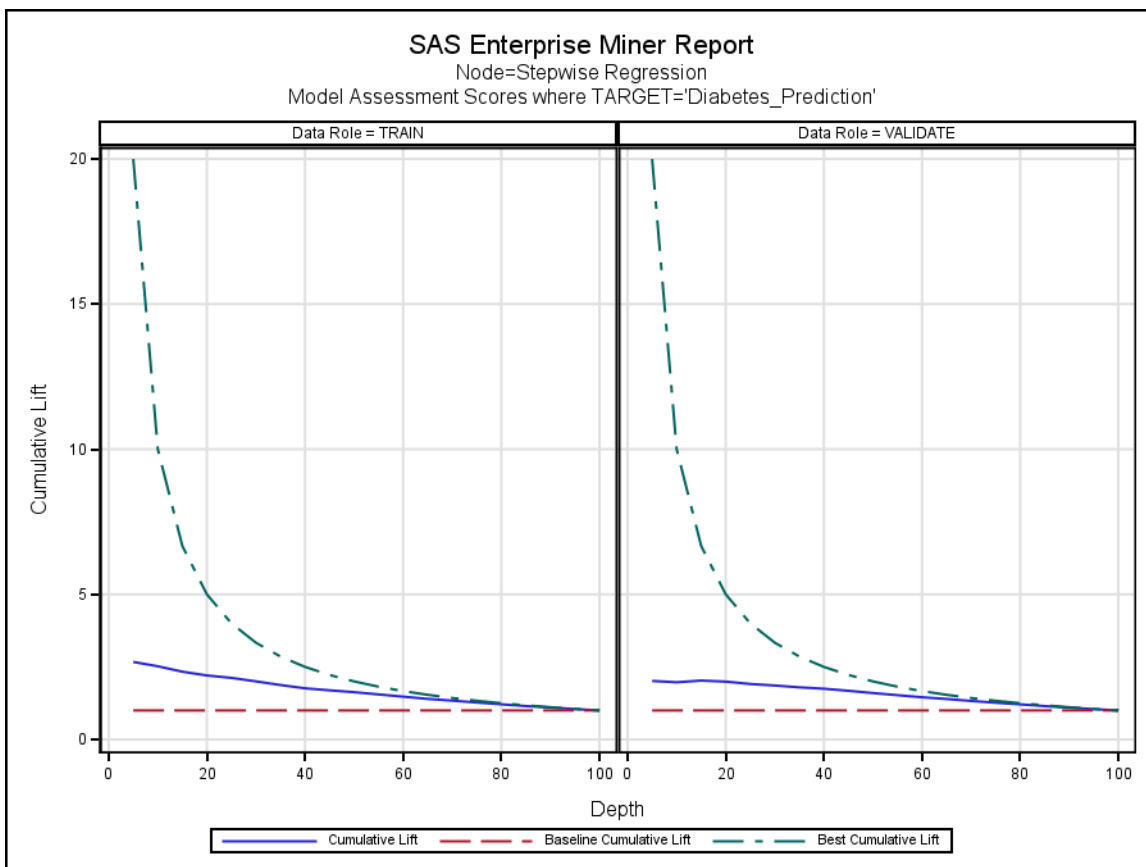
Target=Diabetes_Prediction Target Label=' '

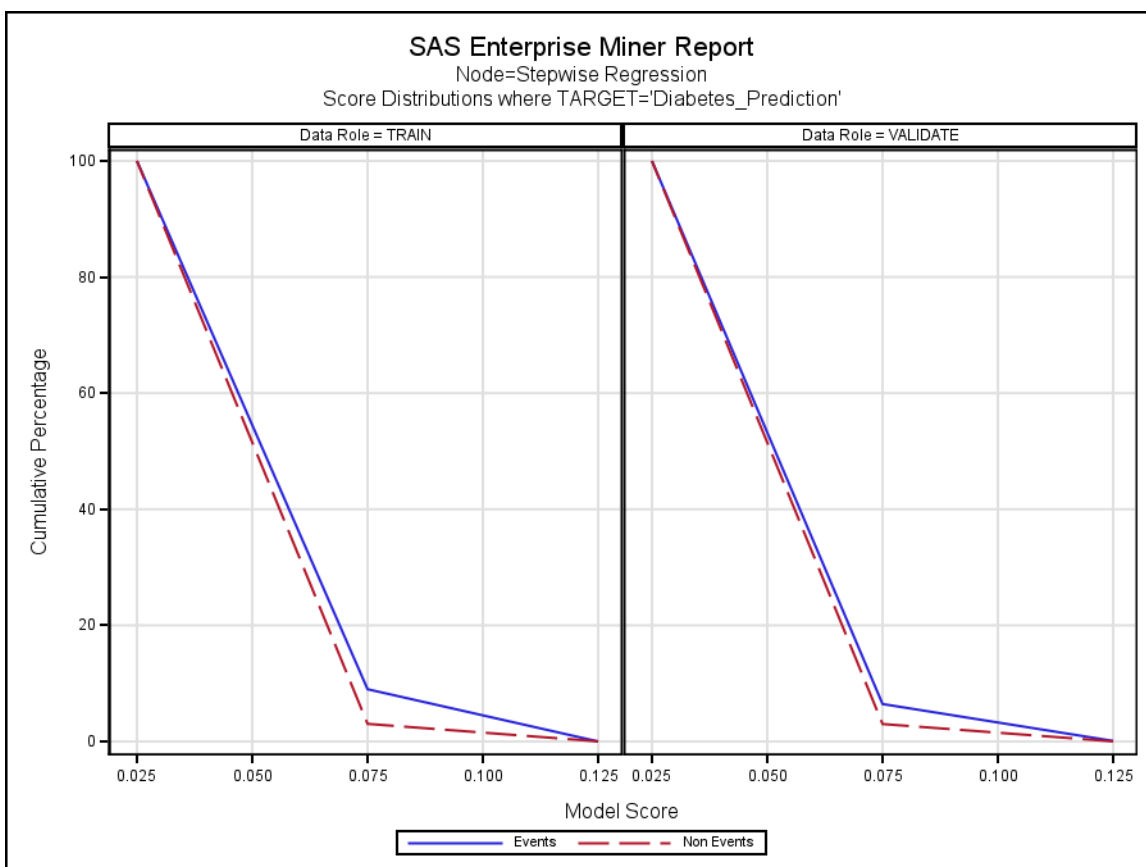
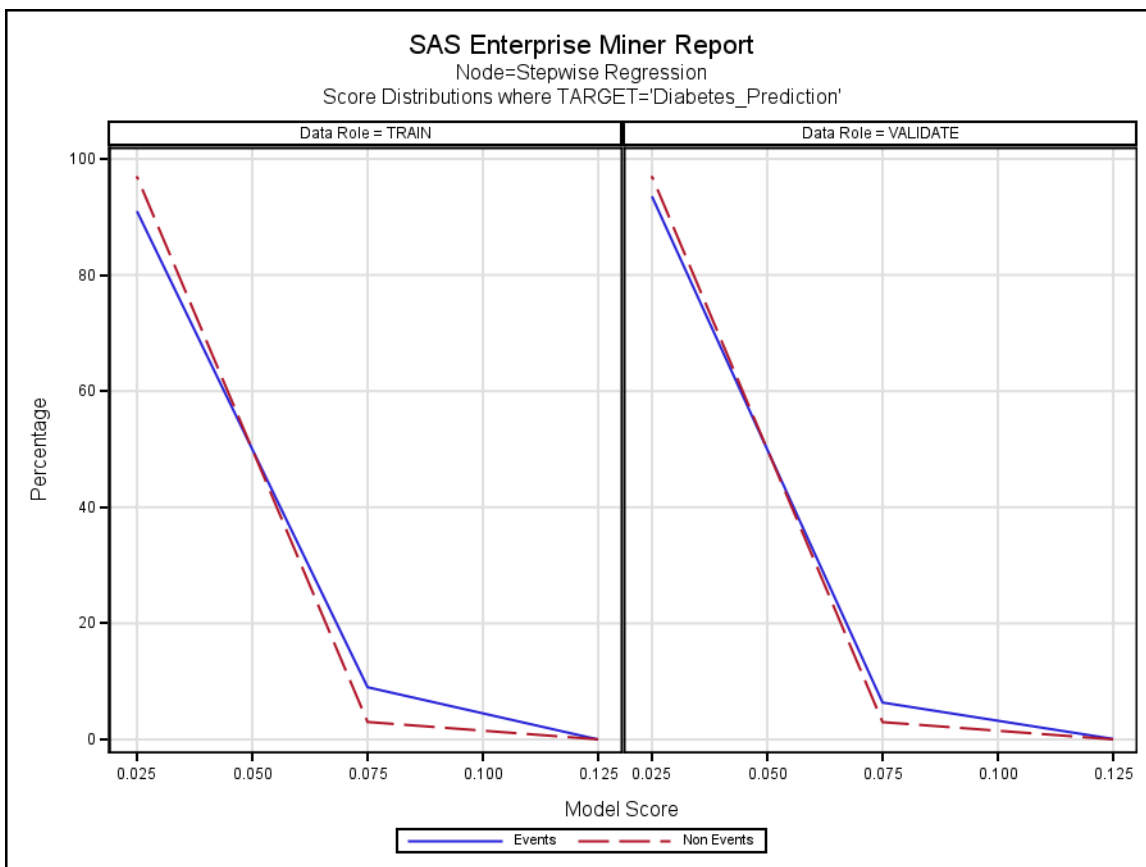
| Label of Statistic | Train | Validation | Test |
|--------------------------------|-----------|------------|-----------|
| Total Degrees of Freedom | 160454.00 | . | . |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Error Function | 63345.86 | 47578.87 | 47439.71 |
| Final Prediction Error | 0.07 | . | . |
| Maximum Absolute Error | 1.00 | 1.00 | 1.00 |
| Mean Square Error | 0.07 | 0.07 | 0.07 |
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Number of Estimate Weights | 56.00 | . | . |
| Root Average Sum of Squares | 0.27 | 0.27 | 0.27 |
| Root Final Prediction Error | 0.27 | . | . |
| Root Mean Squared Error | 0.27 | 0.27 | 0.27 |
| Schwarz's Bayesian Criterion | 64017.07 | . | . |
| Sum of Squared Errors | 17742.41 | 13298.36 | 13290.51 |
| Sum of Case Weights Times Freq | 240681.00 | 180507.00 | 180528.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |

SAS Enterprise Miner Report
Node=Stepwise Regression
Regression Model Effects









Node=Stepwise Regression
Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.10-0.15 | 0 | 0.0000 | 0.0203 | 0.000 | 0.020 |
| 0.05-0.10 | 132 | 8.9980 | 3.0142 | 8.998 | 3.035 |
| 0.00-0.05 | 1335 | 91.0020 | 96.9655 | 100.000 | 100.000 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.10-0.15 | 1 | 0.0908 | 0.0169 | 0.091 | 0.017 |
| 0.05-0.10 | 70 | 6.3579 | 2.9881 | 6.449 | 3.005 |
| 0.00-0.05 | 1030 | 93.5513 | 96.9950 | 100.000 | 100.000 |

SAS Enterprise Miner Report

Node=RegTree B2D6 Summary

Node id = Tree
Node label = RegTree B2D6
Meta path = Ids3 => Part => Tree
Notes =

Node=RegTree B2D6 Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|-------------------|----------------|---------|------------------|-------------|---------|-------------------|------------|---------|
| Component | DecisionTree | | Kass | Y | | Pred | N | |
| AVG | Y | | KassApply | BEFORE | | Predict | Y | |
| AssessMeasure | PROFIT/LOSS | | LeafSize | 5 | | ProfitLoss | NONE | |
| AssessPercentage | 0.25 | | Leafid | Y | | RASE | N | |
| CV | N | | Maxbranch | 2 | | SampleMethod | RANDOM | |
| CVNlter | 10 | | Maxdepth | 6 | | SampleSeed | 12345 | |
| CVRepeat | 1 | | MinCatSize | 5 | | SampleSize | 10000 | |
| CVSeed | 12345 | | MissingValue | USEINSEARCH | | ShowNodeid | Y | |
| ClassColorBy | PERCENTCORRECT | | NSubtree | 1 | | ShowValid | Y | |
| Count | Y | | NodeRole | SEGMENT | | SigLevel | 0.2 | |
| CreateSample | DEFAULT | | NodeSample | 20000 | | SplitPrecision | 4 | |
| Criterion | DEFAULT | | NominalCriterion | PROBCHISQ | | Splitsize | . | |
| Depth | Y | | Nrules | 5 | | Subtree | ASSESSMENT | |
| Dummy | N | | Nsurrs | 0 | | Target | ALL | |
| Exhaustive | 5000 | | NumInputs | 1 | | ToolType | MODEL | |
| Freeze | N | | NumSingleImp | 5 | | TrainMode | BATCH | |
| ImportModel | N | | ObsImportance | N | | UseDecision | N | |
| ImportedTreeData | | | OrdinalCriterion | ENTROPY | | UseMultipleTarget | N | |
| Inputs | N | | PercentCorrect | N | | UsePriors | N | |
| IntColorBy | AVG | | Performance | DISK | | UseVarOnce | N | |
| IntervalCriterion | PROBF | | Precision | 4 | | VarSelection | Y | |

Node=RegTree B2D6 Variable Summary

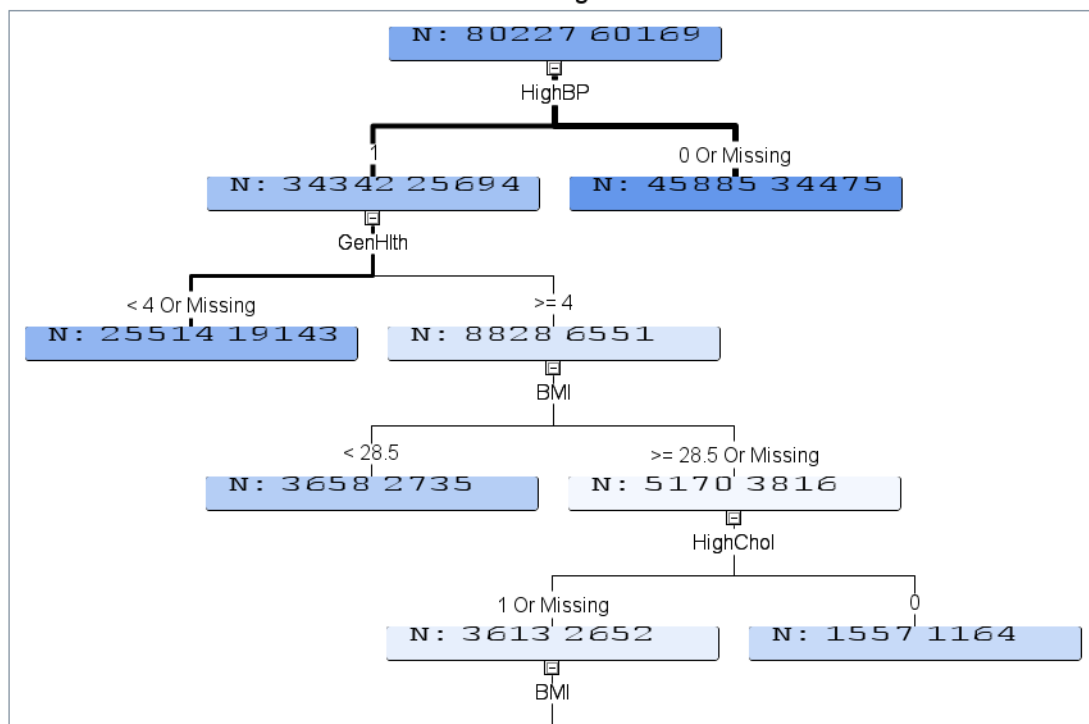
| Role | Level | Frequency Count | Name |
|--------|----------|--------------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |
| ID | INTERVAL | 1 | _dataobs_ |

Node=RegTree B2D6 Model Fit Statistics

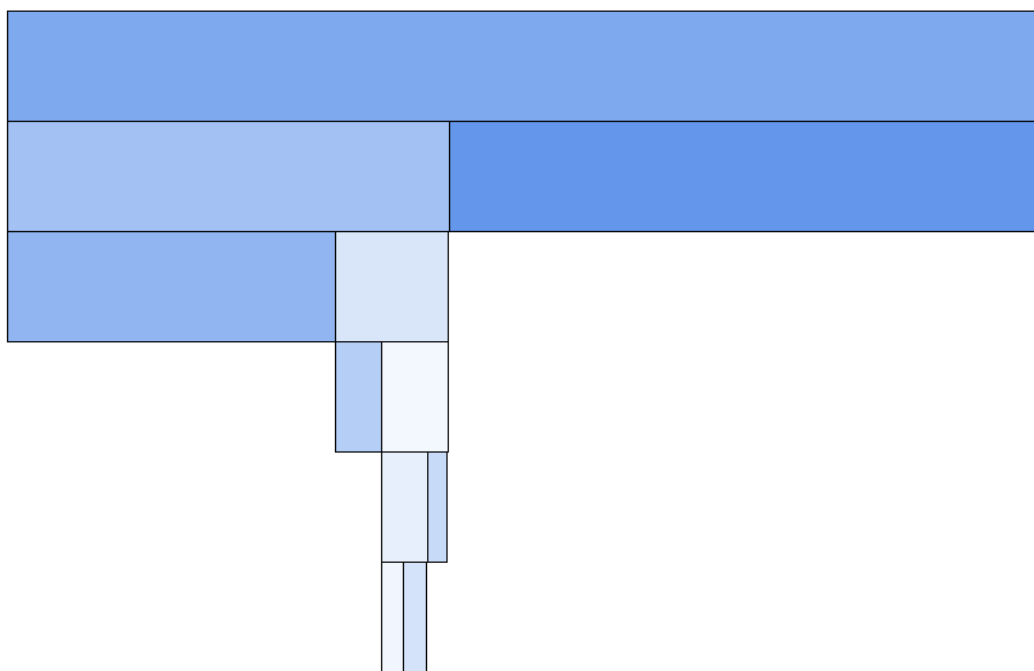
Target=Diabetes_Prediction Target Label='1'

| Label of Statistic | Train | Validation | Test |
|----------------------------|-----------|------------|-----------|
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |
| Maximum Absolute Error | 0.99 | 0.99 | 0.99 |
| Sum of Squared Errors | 19108.87 | 14397.67 | 14381.29 |
| Average Squared Error | 0.08 | 0.08 | 0.08 |
| Root Average Squared Error | 0.28 | 0.28 | 0.28 |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Total Degrees of Freedom | 160454.00 | . | . |

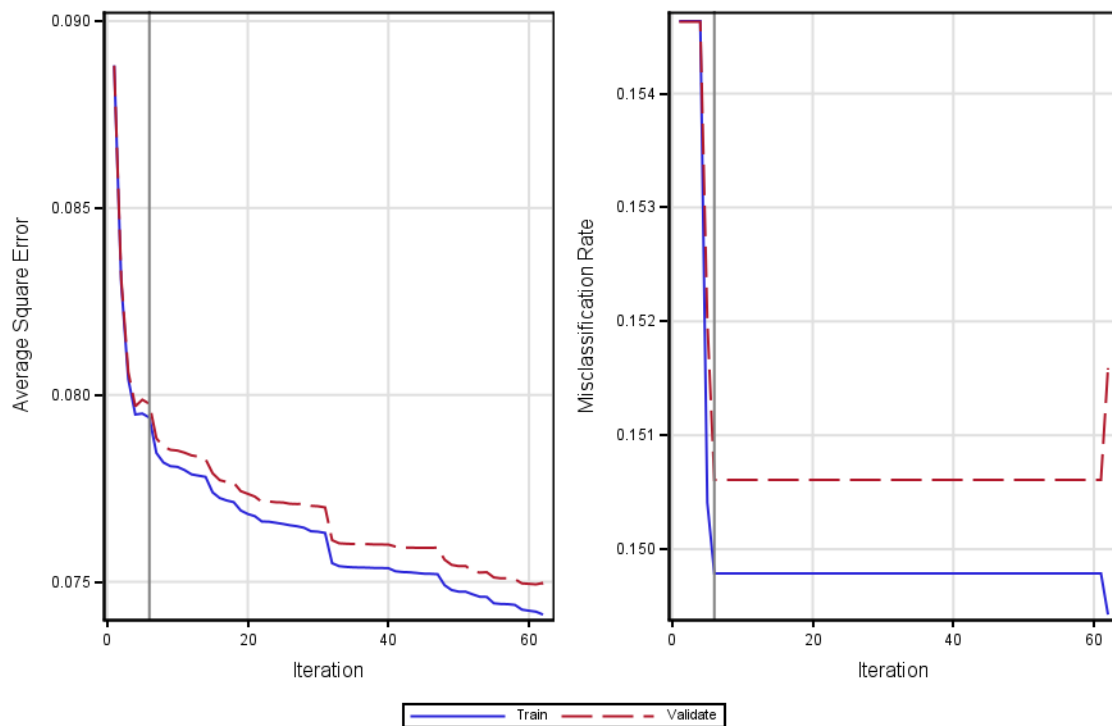
SAS Enterprise Miner Report
Node=RegTree B2D6
Tree Diagram



SAS Enterprise Miner Report
Node=RegTree B2D6
Treemap



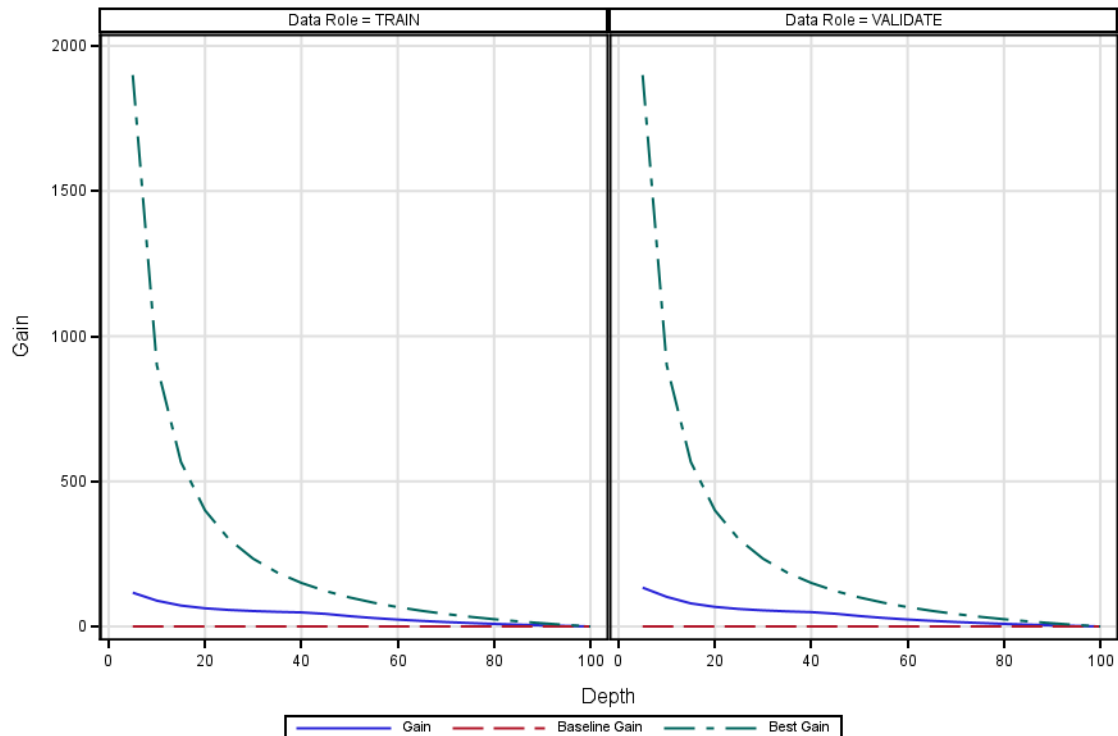
SAS Enterprise Miner Report
Node=RegTree B2D6
Model Iteration Plots



SAS Enterprise Miner Report

Node=RegTree B2D6

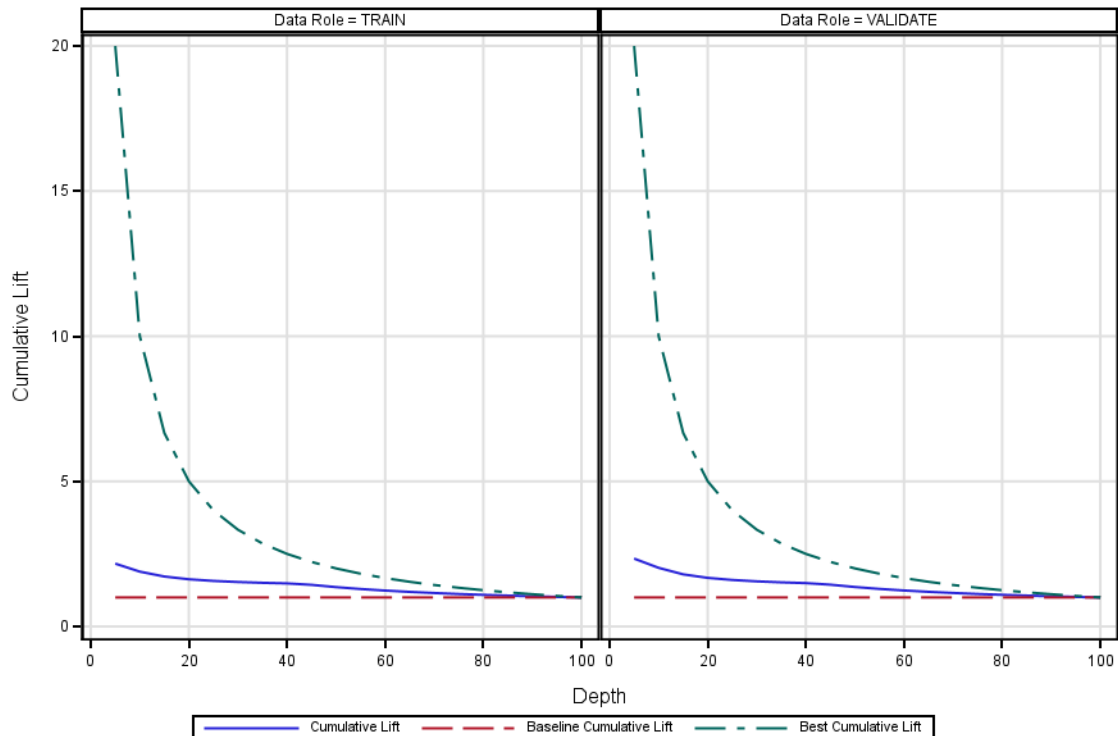
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D6

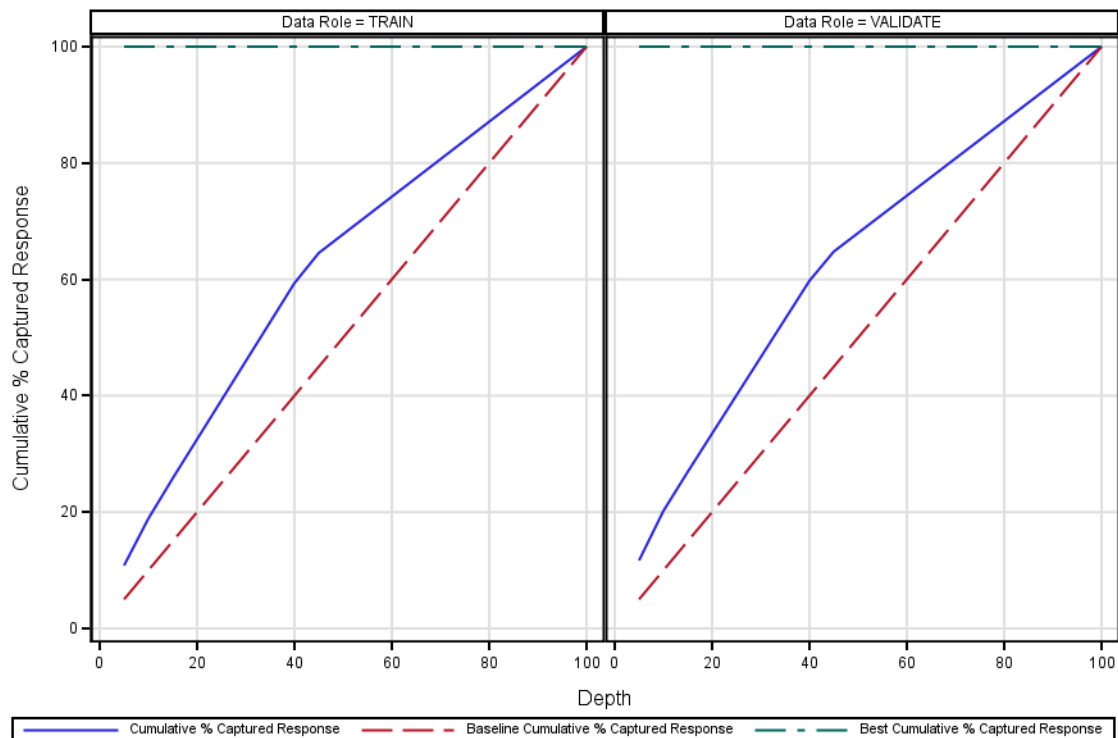
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D6

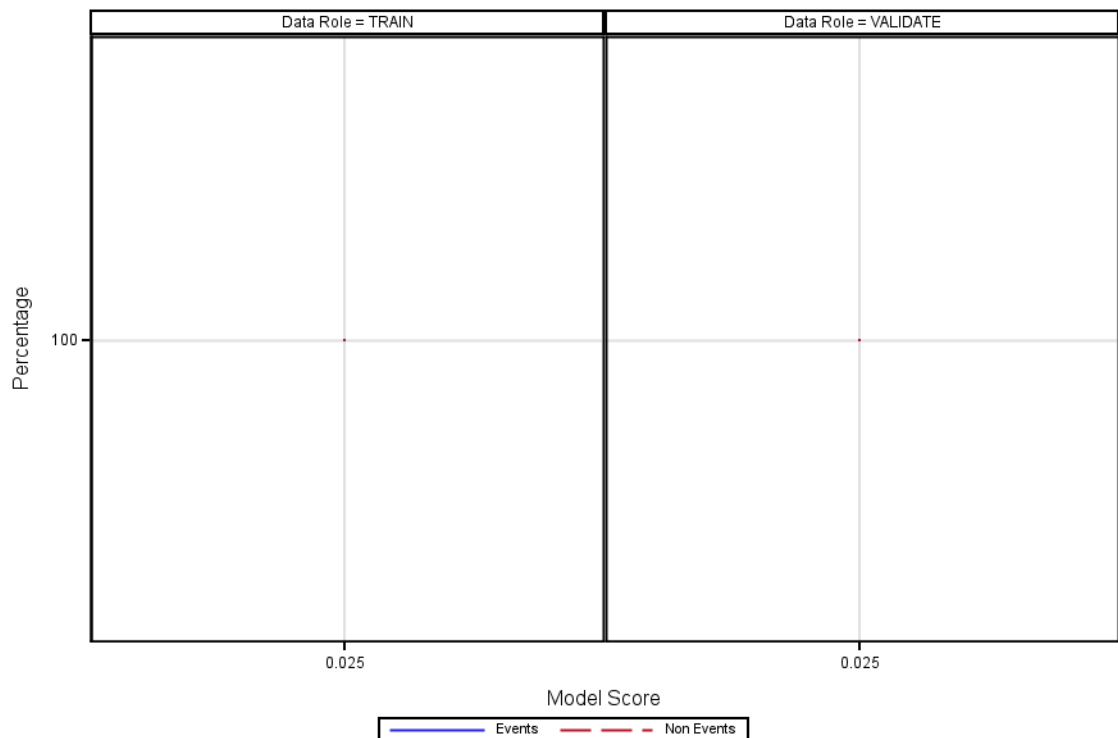
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D6

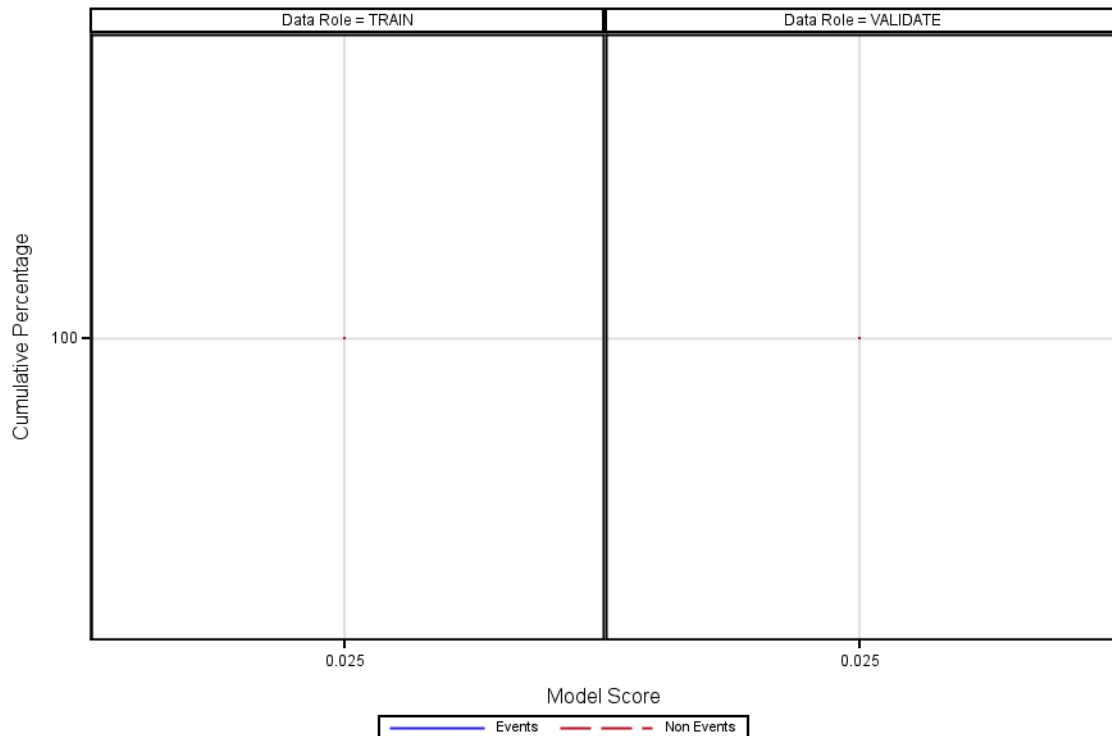
Score Distributions where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D6

Score Distributions where TARGET='Diabetes_Prediction'



Node=RegTree B2D6 Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.00-0.05 | 1467 | 100 | 100 | 100 | 100 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.00-0.05 | 1101 | 100 | 100 | 100 | 100 |

SAS Enterprise Miner Report

Node=RegTree B2D4 Summary

Node id = Tree2
Node label = RegTree B2D4
Meta path = Ids3 => Part => Tree2
Notes =

Node=RegTree B2D4 Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|-------------------|----------------|---------|------------------|-------------|---------|-------------------|------------|---------|
| Component | DecisionTree | | Kass | Y | | Pred | N | |
| AVG | Y | | KassApply | BEFORE | | Predict | Y | |
| AssessMeasure | PROFIT/LOSS | | LeafSize | 5 | | ProfitLoss | NONE | |
| AssessPercentage | 0.25 | | Leafid | Y | | RASE | N | |
| CV | N | | Maxbranch | 2 | | SampleMethod | RANDOM | |
| CVNlter | 10 | | Maxdepth | 4 | 6 | SampleSeed | 12345 | |
| CVRepeat | 1 | | MinCatSize | 5 | | SampleSize | 10000 | |
| CVSeed | 12345 | | MissingValue | USEINSEARCH | | ShowNodeid | Y | |
| ClassColorBy | PERCENTCORRECT | | NSubtree | 1 | | ShowValid | Y | |
| Count | Y | | NodeRole | SEGMENT | | SigLevel | 0.2 | |
| CreateSample | DEFAULT | | NodeSample | 20000 | | SplitPrecision | 4 | |
| Criterion | DEFAULT | | NominalCriterion | PROBCHISQ | | Splitsize | . | |
| Depth | Y | | Nrules | 5 | | Subtree | ASSESSMENT | |
| Dummy | N | | Nsurrs | 0 | | Target | ALL | |
| Exhaustive | 5000 | | NumInputs | 1 | | ToolType | MODEL | |
| Freeze | N | | NumSingleImp | 5 | | TrainMode | BATCH | |
| ImportModel | N | | ObsImportance | N | | UseDecision | N | |
| ImportedTreeData | | | OrdinalCriterion | ENTROPY | | UseMultipleTarget | N | |
| Inputs | N | | PercentCorrect | N | | UsePriors | N | |
| IntColorBy | AVG | | Performance | DISK | | UseVarOnce | N | |
| IntervalCriterion | PROBF | | Precision | 4 | | VarSelection | Y | |

Node=RegTree B2D4 Variable Summary

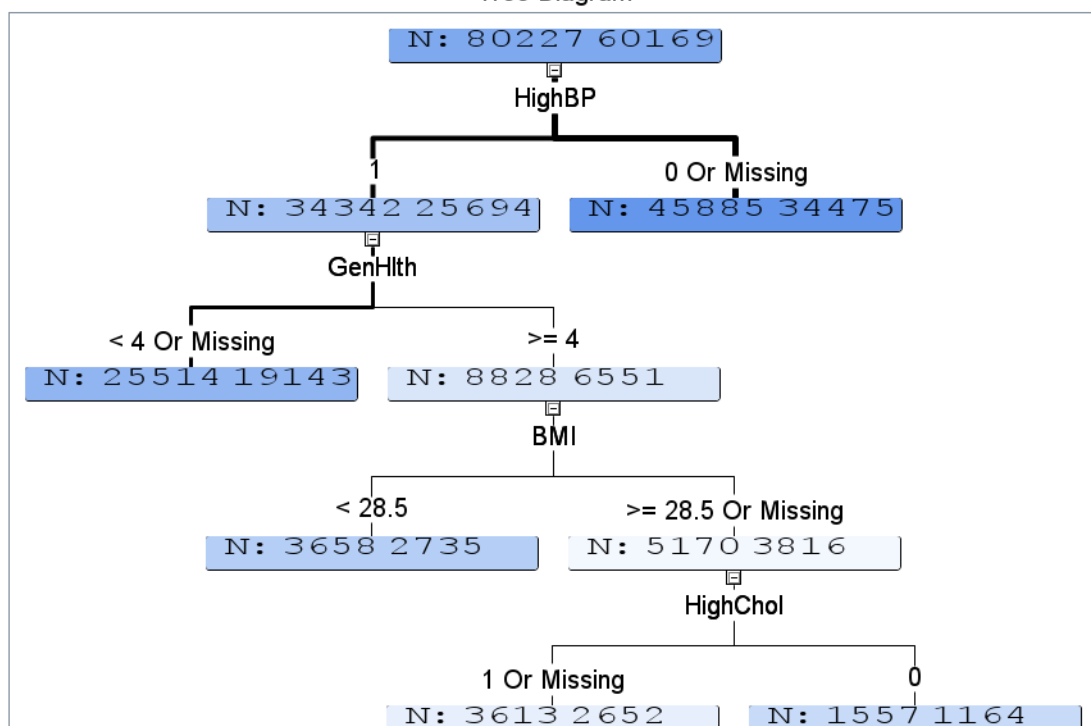
| Role | Level | Frequency Count | Name |
|--------|----------|--------------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |
| ID | INTERVAL | 1 | _dataobs_ |

Node=RegTree B2D4 Model Fit Statistics

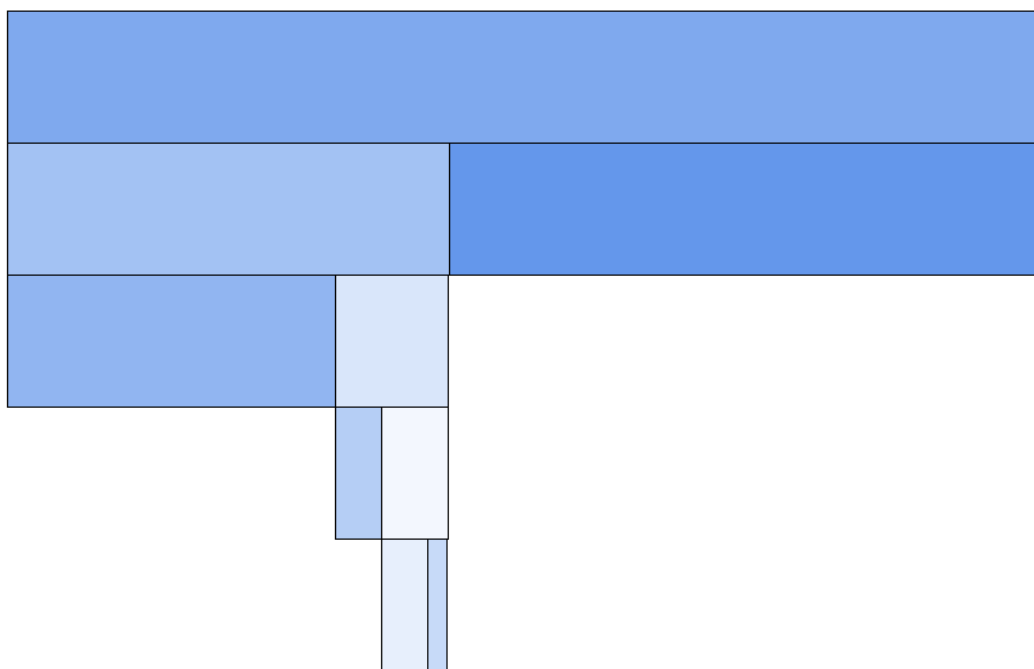
Target=Diabetes_Prediction Target Label=''

| Label of Statistic | Train | Validation | Test |
|----------------------------|-----------|------------|-----------|
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |
| Maximum Absolute Error | 0.99 | 0.99 | 0.99 |
| Sum of Squared Errors | 19134.35 | 14417.52 | 14394.32 |
| Average Squared Error | 0.08 | 0.08 | 0.08 |
| Root Average Squared Error | 0.28 | 0.28 | 0.28 |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Total Degrees of Freedom | 160454.00 | . | . |

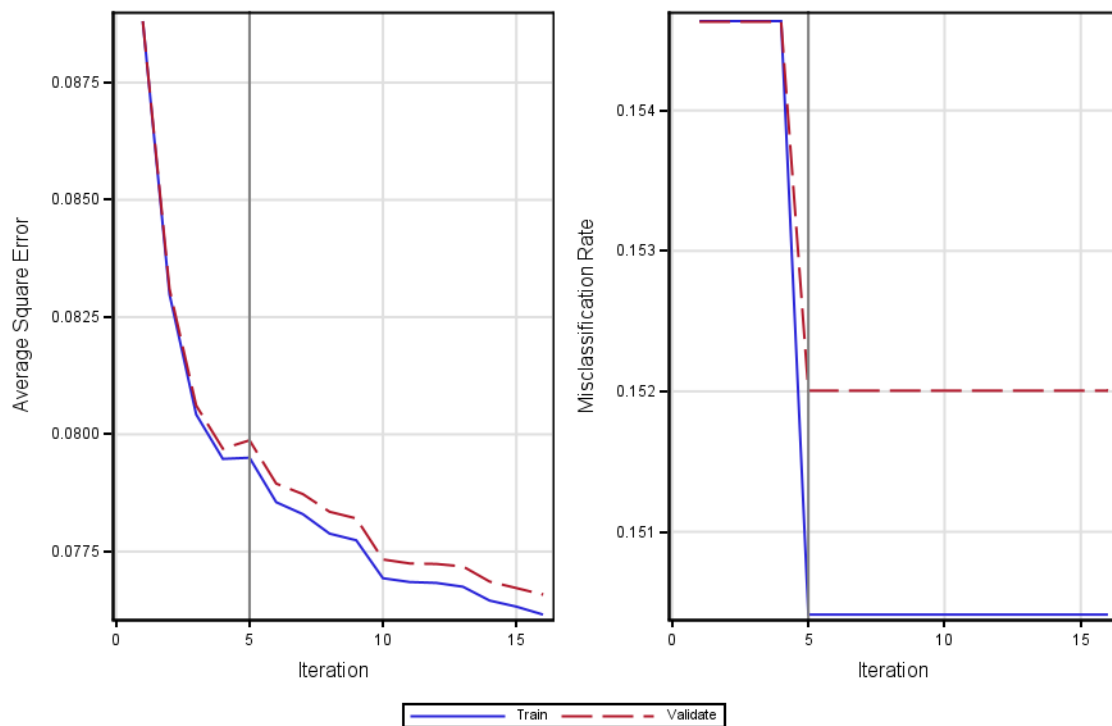
SAS Enterprise Miner Report
Node=RegTree B2D4
Tree Diagram



SAS Enterprise Miner Report
Node=RegTree B2D4
Treemap



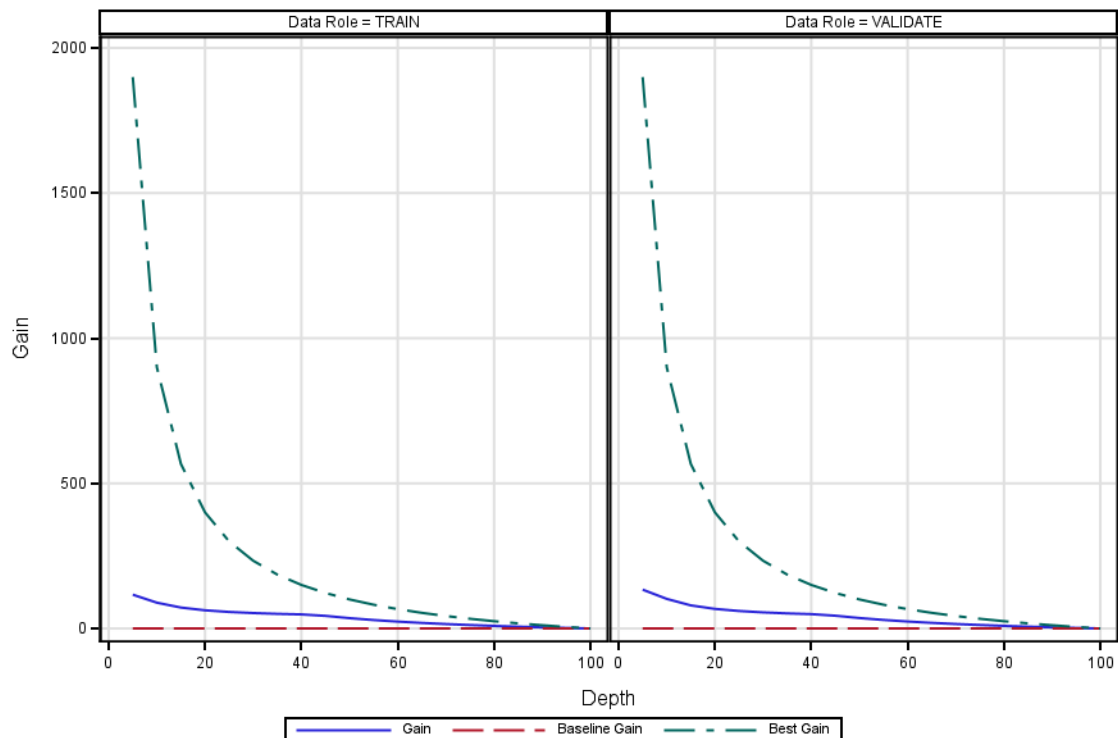
SAS Enterprise Miner Report
Node=RegTree B2D4
Model Iteration Plots



SAS Enterprise Miner Report

Node=RegTree B2D4

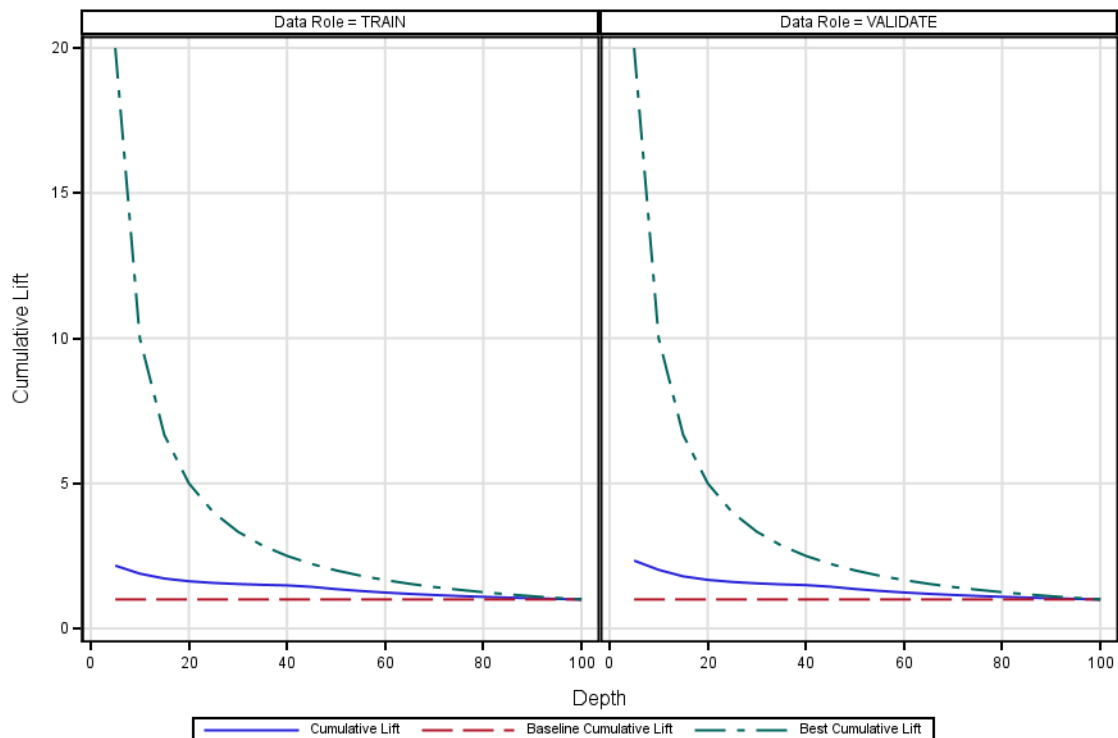
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D4

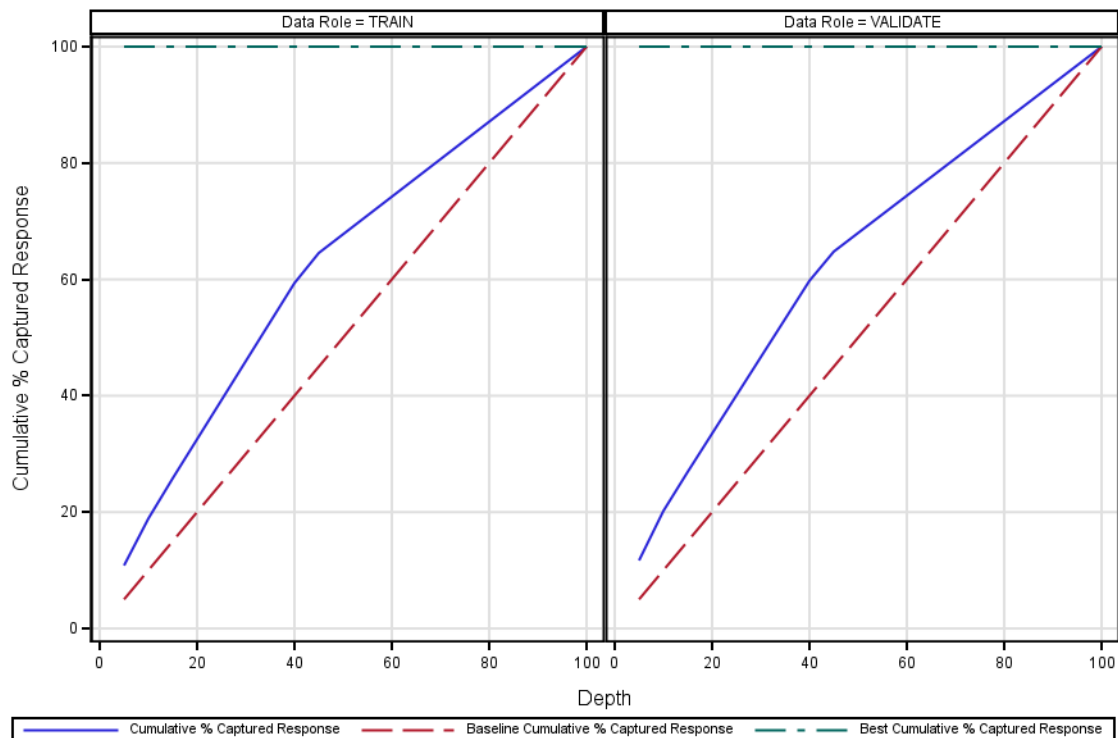
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D4

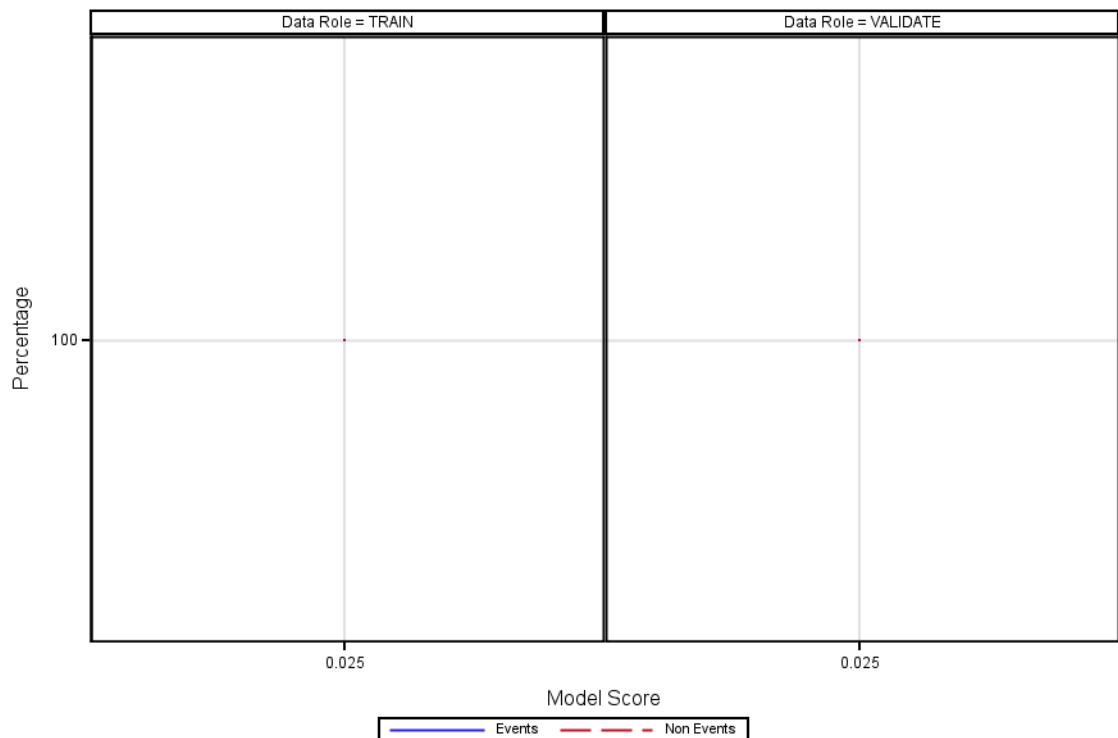
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D4

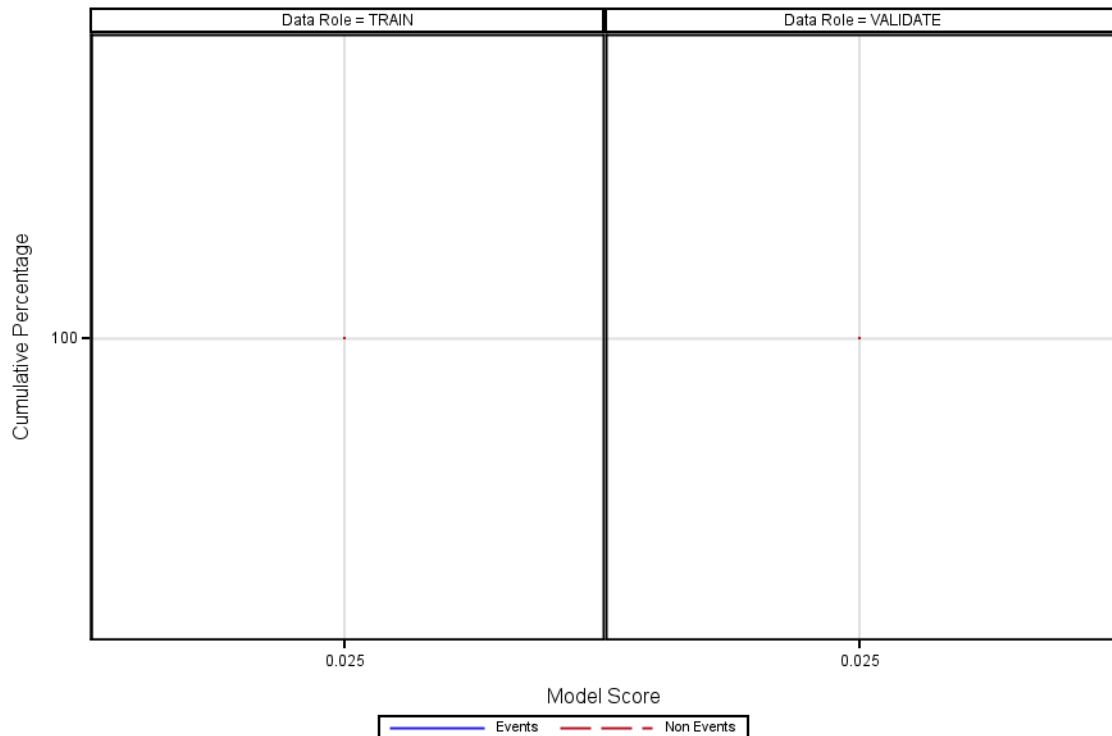
Score Distributions where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D4

Score Distributions where TARGET='Diabetes_Prediction'



Node=RegTree B2D4 Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.00-0.05 | 1467 | 100 | 100 | 100 | 100 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.00-0.05 | 1101 | 100 | 100 | 100 | 100 |

SAS Enterprise Miner Report

Node=RegTree B2D2 Summary

Node id = Tree3
Node label = RegTree B2D2
Meta path = Ids3 => Part => Tree3
Notes =

Node=RegTree B2D2 Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|-------------------|----------------|---------|------------------|-------------|---------|-------------------|------------|---------|
| Component | DecisionTree | | Kass | Y | | Pred | N | |
| AVG | Y | | KassApply | BEFORE | | Predict | Y | |
| AssessMeasure | PROFIT/LOSS | | LeafSize | 5 | | ProfitLoss | NONE | |
| AssessPercentage | 0.25 | | Leafid | Y | | RASE | N | |
| CV | N | | Maxbranch | 2 | | SampleMethod | RANDOM | |
| CVNlter | 10 | | Maxdepth | 2 | 6 | SampleSeed | 12345 | |
| CVRepeat | 1 | | MinCatSize | 5 | | SampleSize | 10000 | |
| CVSeed | 12345 | | MissingValue | USEINSEARCH | | ShowNodeid | Y | |
| ClassColorBy | PERCENTCORRECT | | NSubtree | 1 | | ShowValid | Y | |
| Count | Y | | NodeRole | SEGMENT | | SigLevel | 0.2 | |
| CreateSample | DEFAULT | | NodeSample | 20000 | | SplitPrecision | 4 | |
| Criterion | DEFAULT | | NominalCriterion | PROBCHISQ | | Splitsize | . | |
| Depth | Y | | Nrules | 5 | | Subtree | ASSESSMENT | |
| Dummy | N | | Nsurrs | 0 | | Target | ALL | |
| Exhaustive | 5000 | | NumInputs | 1 | | ToolType | MODEL | |
| Freeze | N | | NumSingleImp | 5 | | TrainMode | BATCH | |
| ImportModel | N | | ObsImportance | N | | UseDecision | N | |
| ImportedTreeData | | | OrdinalCriterion | ENTROPY | | UseMultipleTarget | N | |
| Inputs | N | | PercentCorrect | N | | UsePriors | N | |
| IntColorBy | AVG | | Performance | DISK | | UseVarOnce | N | |
| IntervalCriterion | PROBF | | Precision | 4 | | VarSelection | Y | |

Node=RegTree B2D2 Variable Summary

| Role | Level | Frequency Count | Name |
|--------|----------|-----------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |
| ID | INTERVAL | 1 | _dataobs_ |

Node=RegTree B2D2 Model Fit Statistics

Target=Diabetes_Prediction Target Label=' '

| Label of Statistic | Train | Validation | Test |
|----------------------------|-----------|------------|-----------|
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |
| Maximum Absolute Error | 0.98 | 0.98 | 0.98 |
| Sum of Squared Errors | 21375.22 | 16030.83 | 16038.69 |
| Average Squared Error | 0.09 | 0.09 | 0.09 |
| Root Average Squared Error | 0.30 | 0.30 | 0.30 |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Total Degrees of Freedom | 160454.00 | . | . |

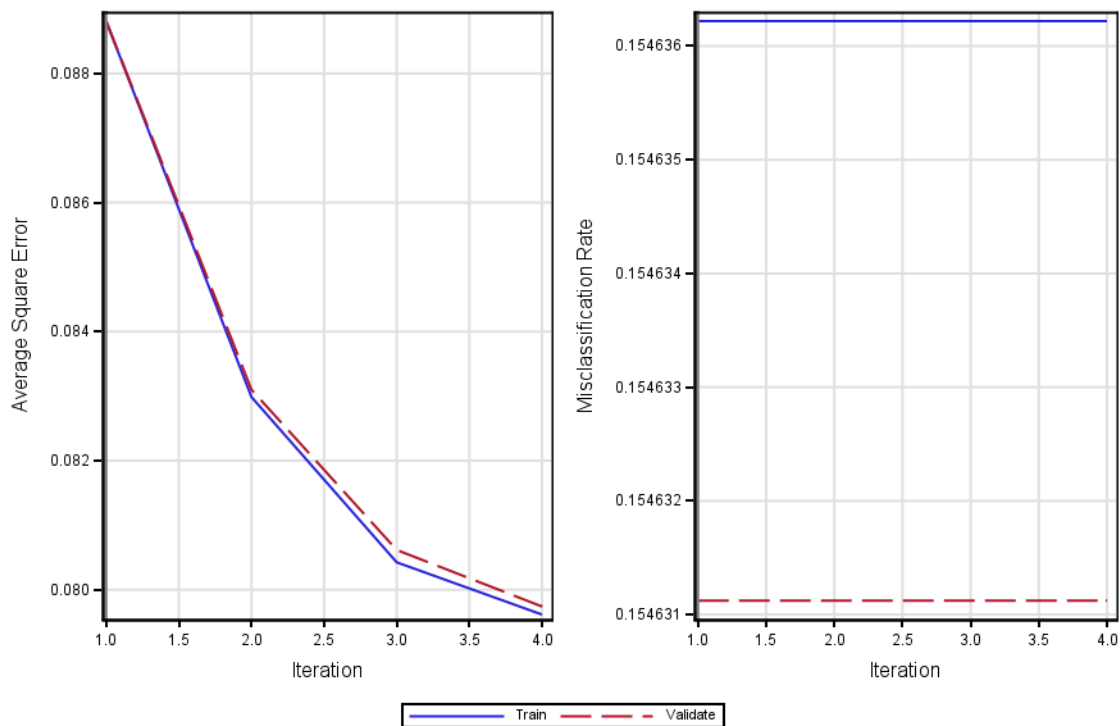
SAS Enterprise Miner Report
Node=RegTree B2D2
Tree Diagram

N: 80227 60169

SAS Enterprise Miner Report
Node=RegTree B2D2
Treemap



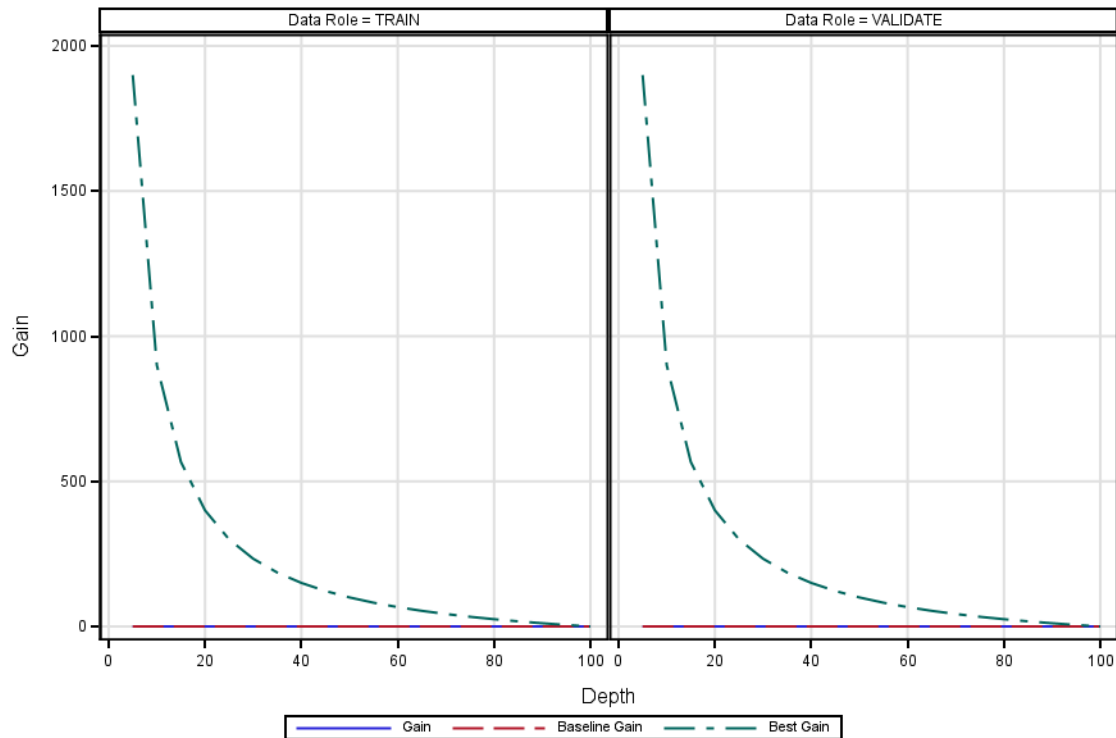
SAS Enterprise Miner Report
Node=RegTree B2D2
Model Iteration Plots



SAS Enterprise Miner Report

Node=RegTree B2D2

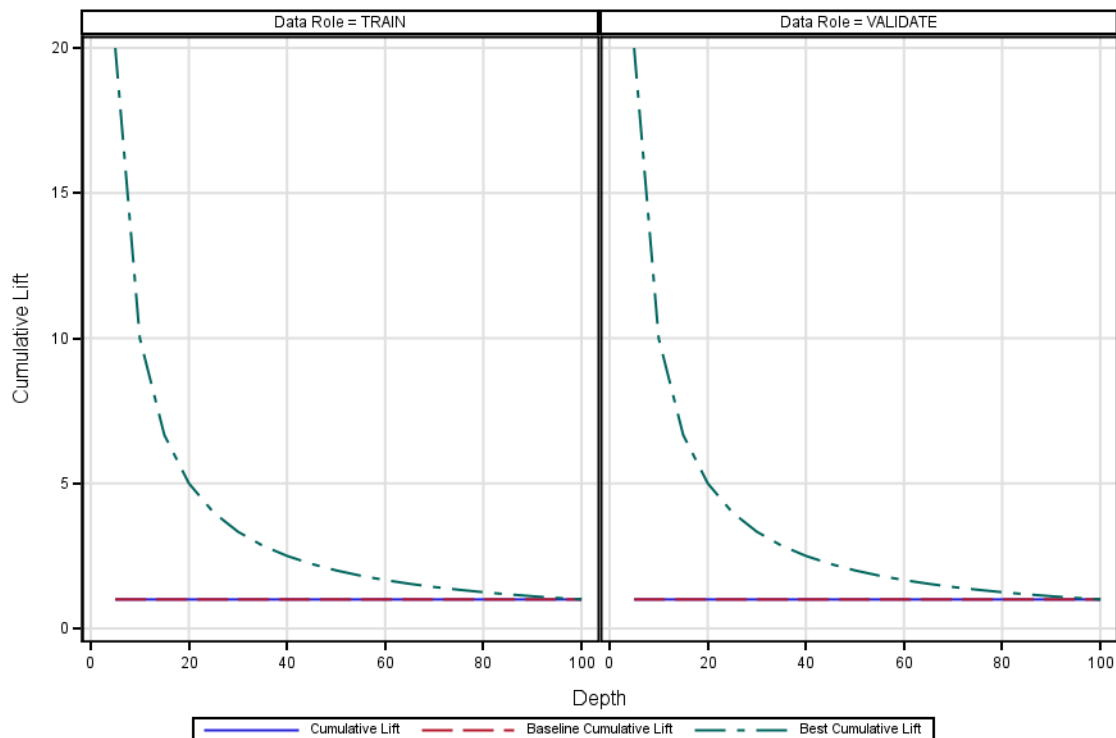
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D2

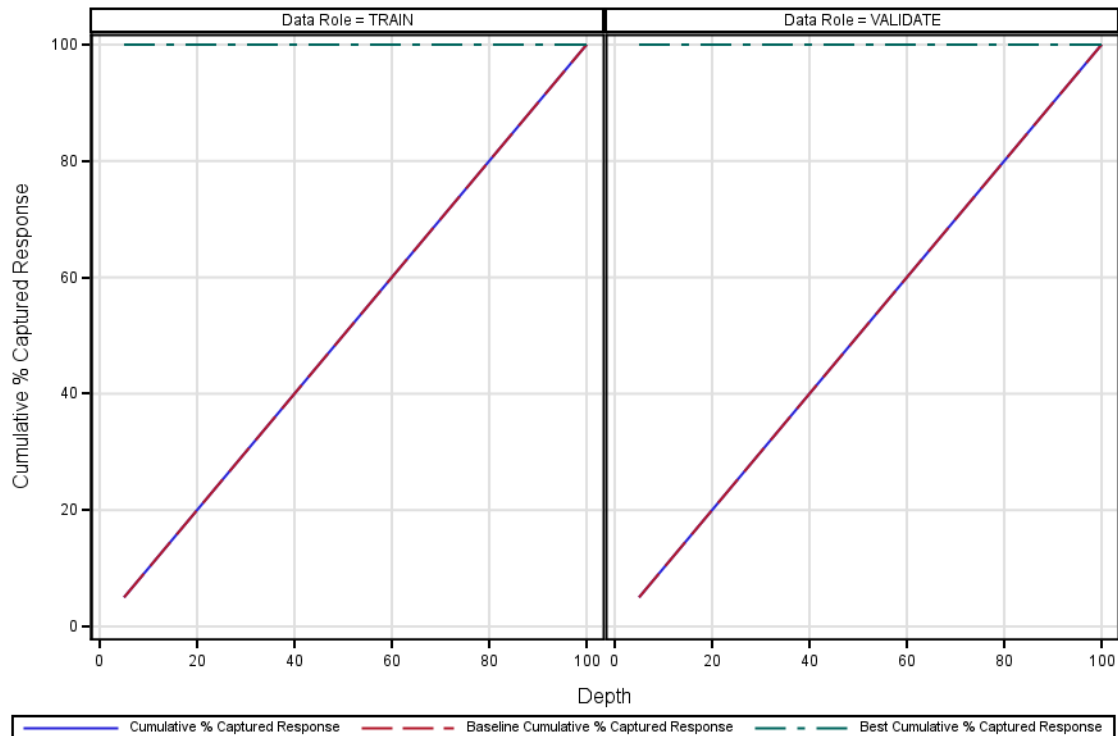
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B2D2

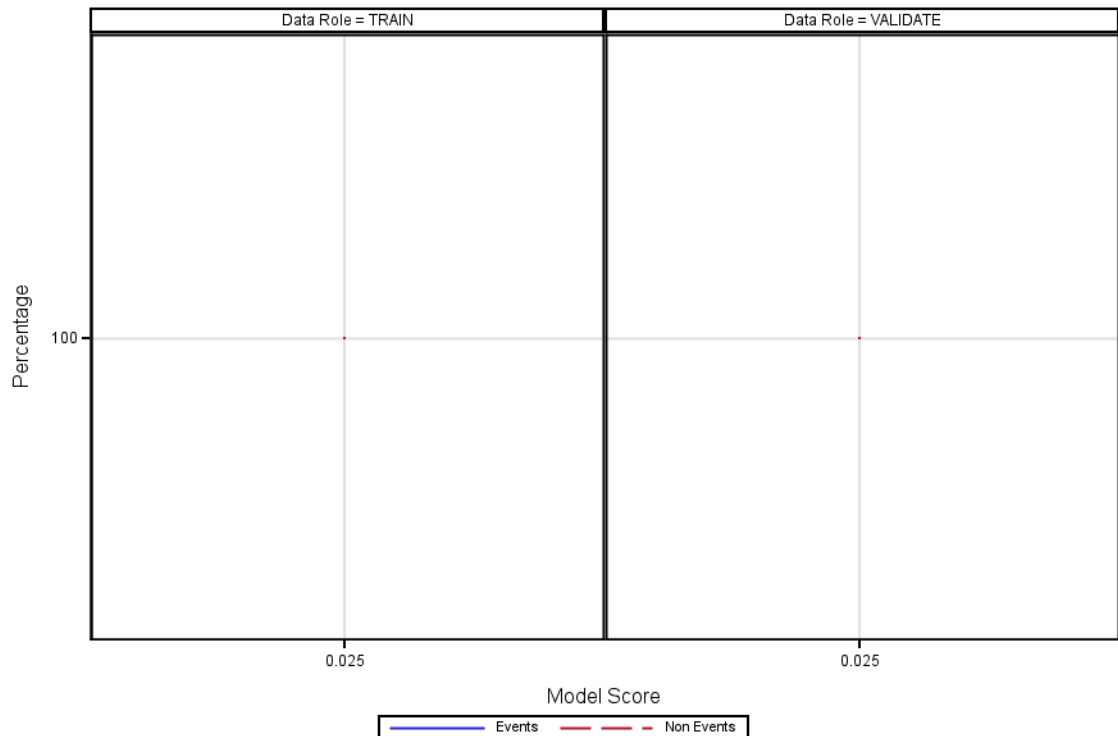
Model Assessment Scores where TARGET='Diabetes_Prediction'

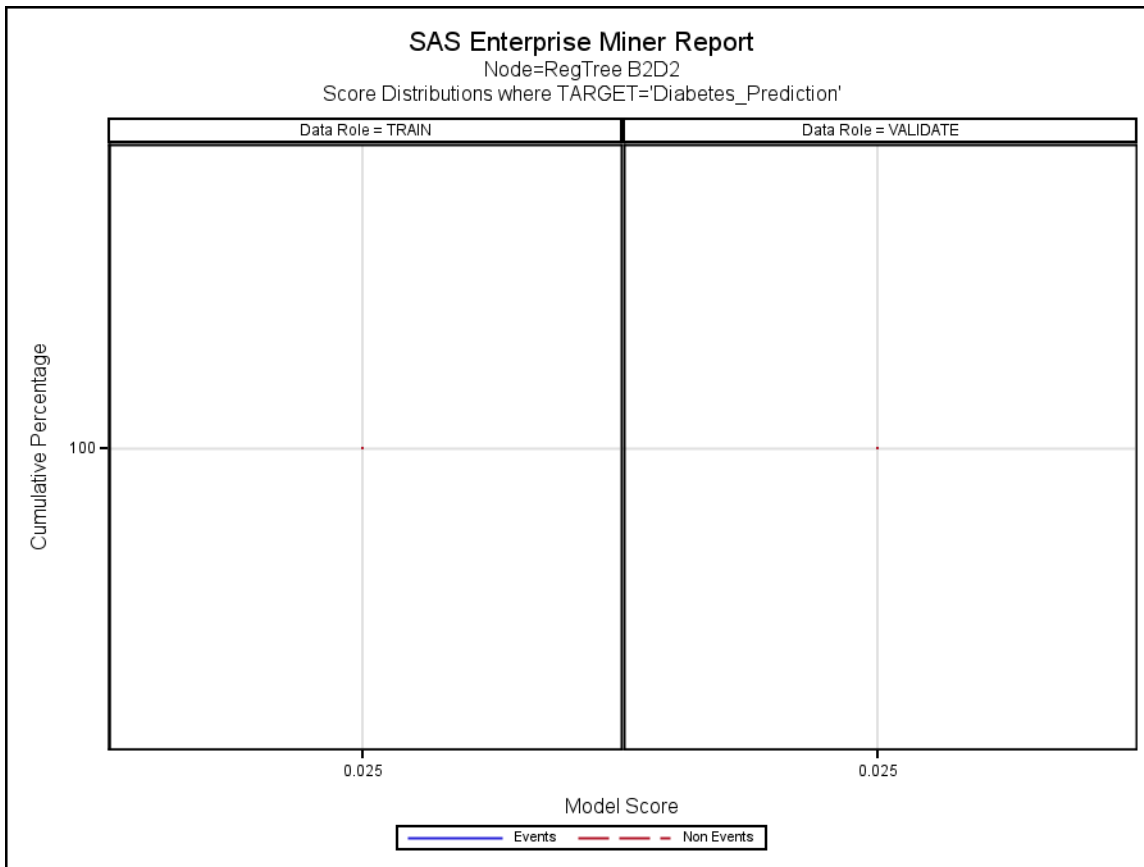


SAS Enterprise Miner Report

Node=RegTree B2D2

Score Distributions where TARGET='Diabetes_Prediction'





Node=RegTree B2D2 Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.00-0.05 | 1467 | 100 | 100 | 100 | 100 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.00-0.05 | 1101 | 100 | 100 | 100 | 100 |

SAS Enterprise Miner Report

Node=RegTree B3D6 Summary

Node id = Tree4
Node label = RegTree B3D6
Meta path = Ids3 => Part => Tree4
Notes =

Node=RegTree B3D6 Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|-------------------|----------------|---------|------------------|-------------|---------|-------------------|------------|---------|
| Component | DecisionTree | | Kass | Y | | Pred | N | |
| AVG | Y | | KassApply | BEFORE | | Predict | Y | |
| AssessMeasure | PROFIT/LOSS | | LeafSize | 5 | | ProfitLoss | NONE | |
| AssessPercentage | 0.25 | | Leafid | Y | | RASE | N | |
| CV | N | | Maxbranch | 3 | 2 | SampleMethod | RANDOM | |
| CVNIter | 10 | | Maxdepth | 6 | | SampleSeed | 12345 | |
| CVRepeat | 1 | | MinCatSize | 5 | | SampleSize | 10000 | |
| CVSeed | 12345 | | MissingValue | USEINSEARCH | | ShowNodeId | Y | |
| ClassColorBy | PERCENTCORRECT | | NSubtree | 1 | | ShowValid | Y | |
| Count | Y | | NodeRole | SEGMENT | | SigLevel | 0.2 | |
| CreateSample | DEFAULT | | NodeSample | 20000 | | SplitPrecision | 4 | |
| Criterion | DEFAULT | | NominalCriterion | PROBCHISQ | | Splitsize | . | |
| Depth | Y | | Nrules | 5 | | Subtree | ASSESSMENT | |
| Dummy | N | | Nsurrs | 0 | | Target | ALL | |
| Exhaustive | 5000 | | NumInputs | 1 | | ToolType | MODEL | |
| Freeze | N | | NumSingleImp | 5 | | TrainMode | BATCH | |
| ImportModel | N | | ObsImportance | N | | UseDecision | N | |
| ImportedTreeData | | | OrdinalCriterion | ENTROPY | | UseMultipleTarget | N | |
| Inputs | N | | PercentCorrect | N | | UsePriors | N | |
| IntColorBy | AVG | | Performance | DISK | | UseVarOnce | N | |
| IntervalCriterion | PROBF | | Precision | 4 | | VarSelection | Y | |

Node=RegTree B3D6 Variable Summary

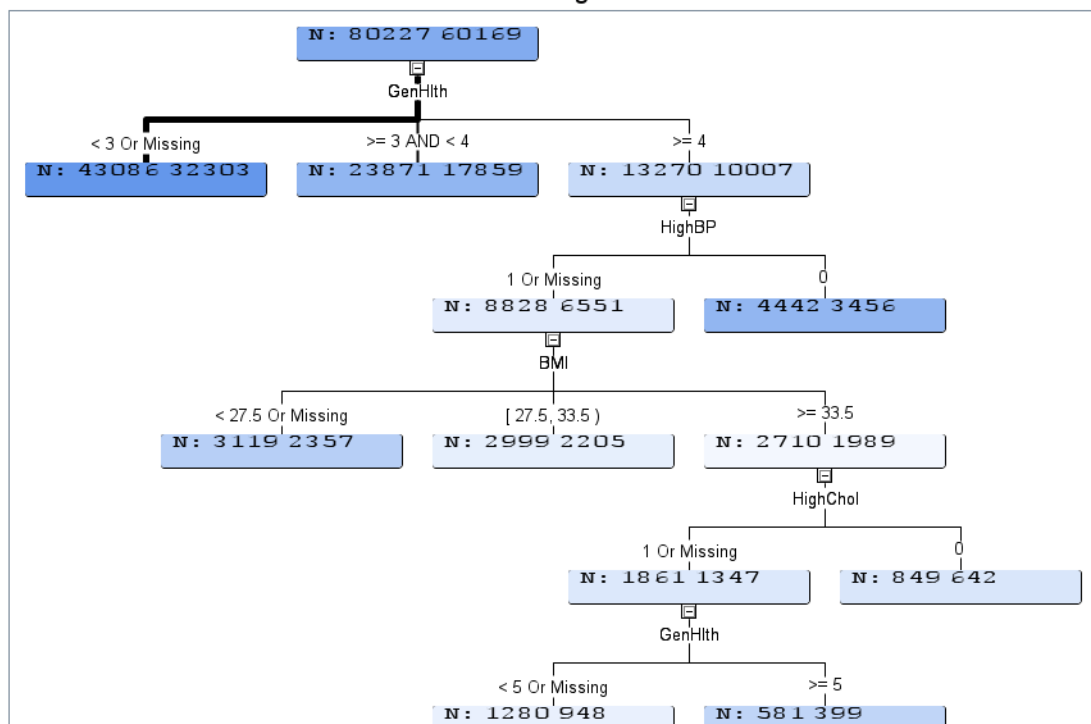
| Role | Level | Frequency Count | Name |
|--------|----------|--------------------|---|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| INPUT | BINARY | 14 | AnyHealthcare CholCheck DiffWalk Fruits HeartDiseaseorAttack HighBP HighChol HvyAlcoholConsump NoDocbcCost PhysActivity Sex Smoker Stroke Veggies |
| INPUT | INTERVAL | 2 | Age BMI |
| INPUT | ORDINAL | 5 | Education GenHlth Income MentHlth PhysHlth |
| ID | INTERVAL | 1 | _dataobs_ |

Node=RegTree B3D6 Model Fit Statistics

Target=Diabetes_Prediction Target Label=''

| Label of Statistic | Train | Validation | Test |
|----------------------------|-----------|------------|-----------|
| Sum of Frequencies | 80227.00 | 60169.00 | 60176.00 |
| Misclassification Rate | 0.15 | 0.15 | 0.15 |
| Maximum Absolute Error | 0.99 | 0.99 | 0.99 |
| Sum of Squared Errors | 19150.93 | 14418.33 | 14374.08 |
| Average Squared Error | 0.08 | 0.08 | 0.08 |
| Root Average Squared Error | 0.28 | 0.28 | 0.28 |
| Divisor for ASE | 240681.00 | 180507.00 | 180528.00 |
| Total Degrees of Freedom | 160454.00 | . | . |

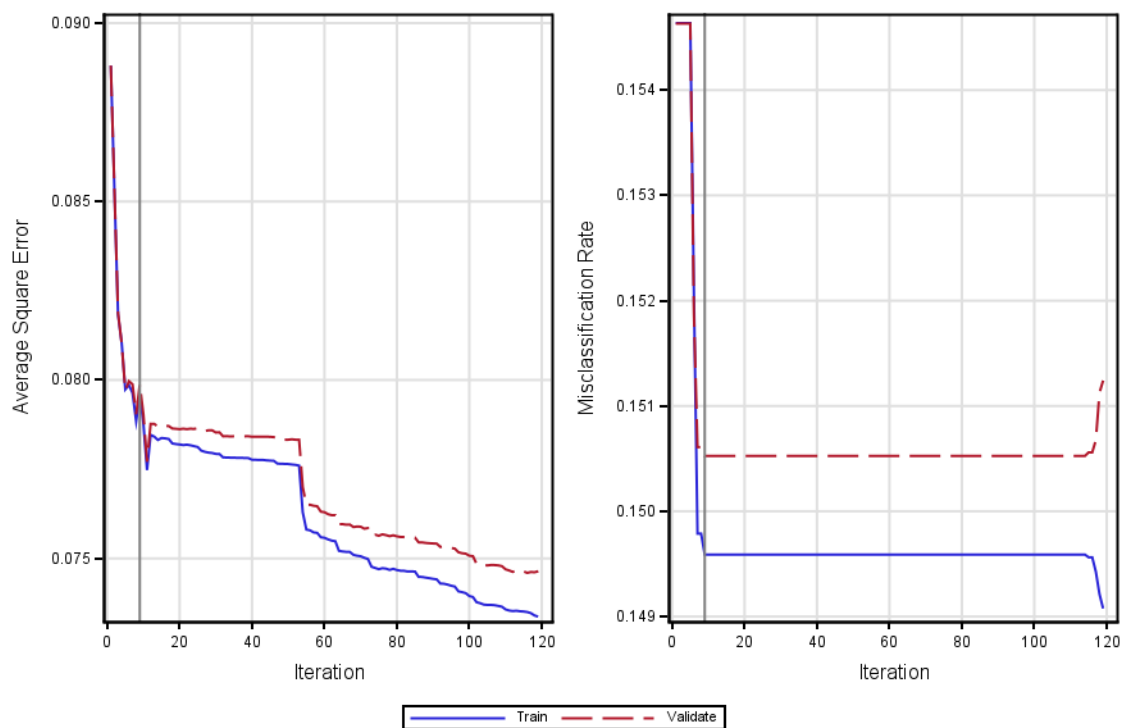
SAS Enterprise Miner Report
Node=RegTree B3D6
Tree Diagram



SAS Enterprise Miner Report
Node=RegTree B3D6
Treemap



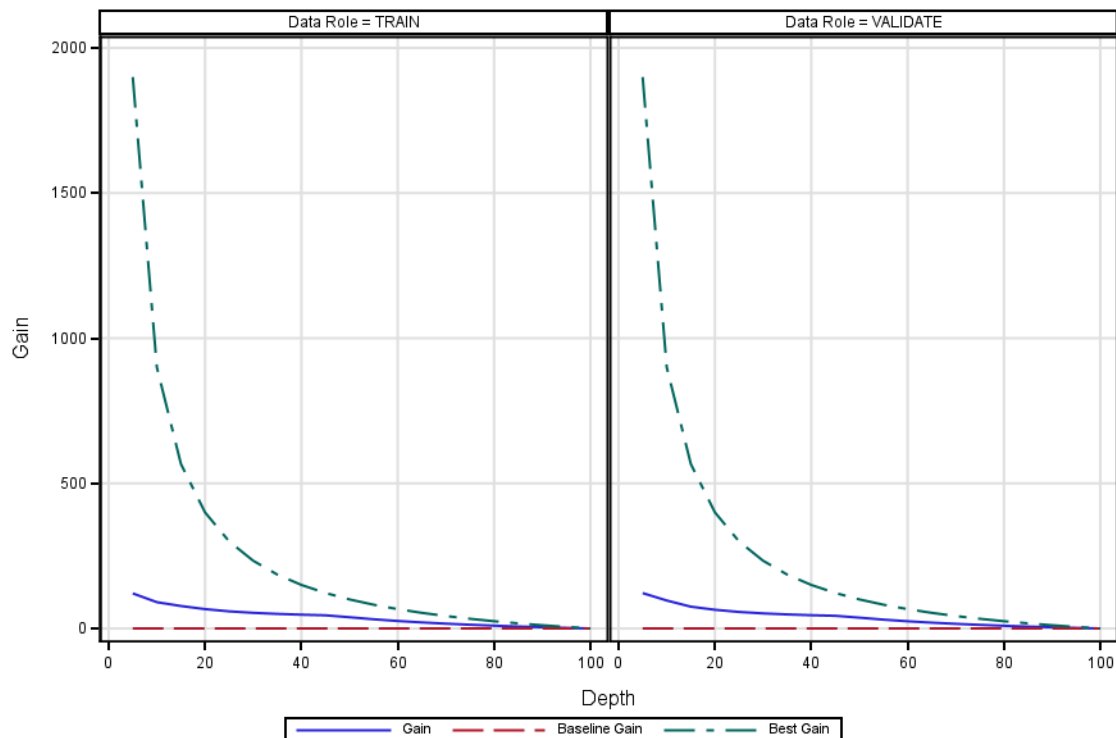
SAS Enterprise Miner Report
Node=RegTree B3D6
Model Iteration Plots



SAS Enterprise Miner Report

Node=RegTree B3D6

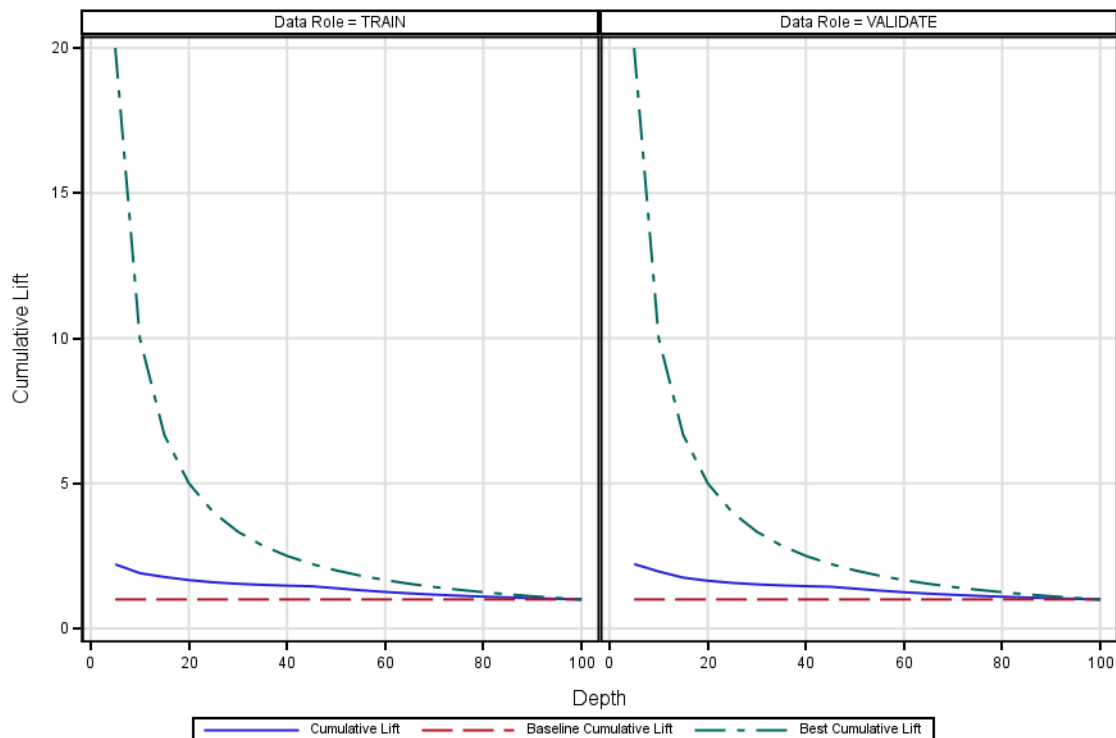
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B3D6

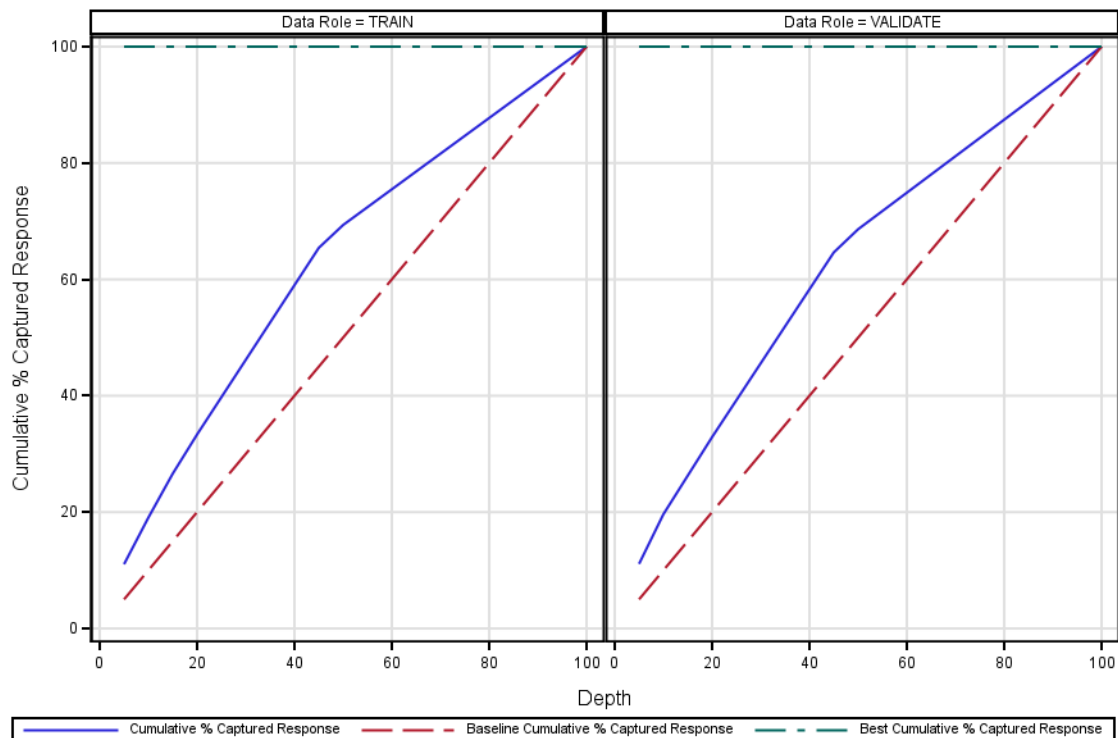
Model Assessment Scores where TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree B3D6

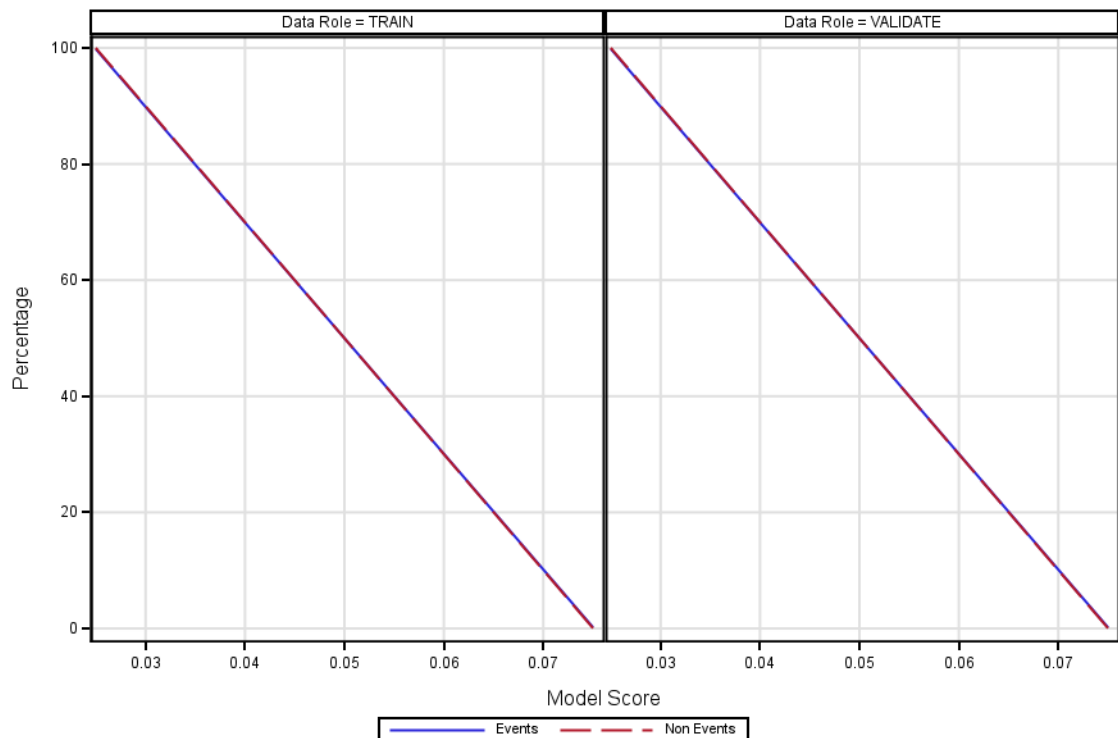
Model Assessment Scores where TARGET='Diabetes_Prediction'

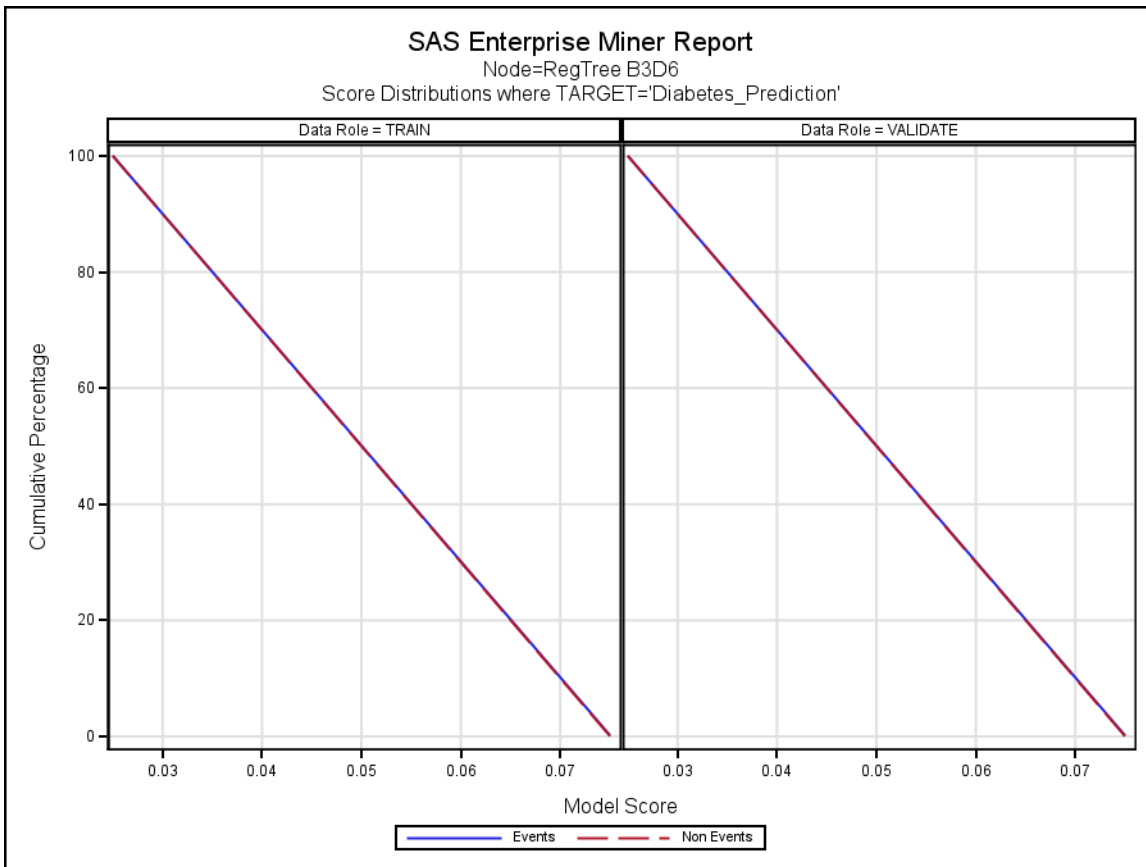


SAS Enterprise Miner Report

Node=RegTree B3D6

Score Distributions where TARGET='Diabetes_Prediction'





Node=RegTree B3D6 Score Distributions

Target Variable=Diabetes_Prediction Data Role=TRAIN

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.05-0.10 | 3 | 0.2045 | 0.0457 | 0.204 | 0.046 |
| 0.00-0.05 | 1464 | 99.7955 | 99.9543 | 100.000 | 100.000 |

Target Variable=Diabetes_Prediction Data Role=VALIDATE

| Posterior Probability Range | Number of Events | Percentage of Events | Percentage of Nonevents | Cumulative Percentage of Events | Cumulative Percentage of Nonevents |
|-----------------------------|------------------|----------------------|-------------------------|---------------------------------|------------------------------------|
| 0.05-0.10 | 2 | 0.1817 | 0.0389 | 0.182 | 0.039 |
| 0.00-0.05 | 1099 | 99.8183 | 99.9611 | 100.000 | 100.000 |

SAS Enterprise Miner Report

Node=MLR ModelComparison Summary

Node id = MdlComp2
Node label = MLR ModelComparison
Meta path = Ids3 => Part => Reg2 => MdlComp2
Notes =

Node=MLR ModelComparison Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|-----------------------------|-------------------------------|---------|------------------------|---------|---------|----------------------|---------------------|---------|
| Component | ModelCompare | | NumberOfReportedLevels | 1E-6 | | SelectionData | DEFAULT | |
| AssessAllTargetLevels | N | | NumberOfBins | 20 | | SelectionDepth | 10 | |
| DecileBin | 20 | | ProfitEpsilon | 1E-6 | | SelectionTable | TRAIN | TABLE |
| HPCriteria | DEFAULT | | RecomputeAssess | N | | StatisticUsed | _VMISC_ | |
| LiftEpsilon | 1E-6 | | RocChart | Y | | TargetLabel | | |
| ModelCriteria | Valid: Misclassification Rate | | RocEpsilon | 0.01 | | TargetName | Diabetes_Prediction | |
| ModelDescription | Forward Regression | | RoiEpsilon | 1E-6 | | classViyaCriteria | DEFAULT | |
| ModelId | Reg2 | | ScoreDistBin | 20 | | intervalViyaCriteria | DEFAULT | |
| NormalizeReportingVariables | Y | | SelectionCriteria | DEFAULT | | | | |

Node=MLR ModelComparison Variable Summary

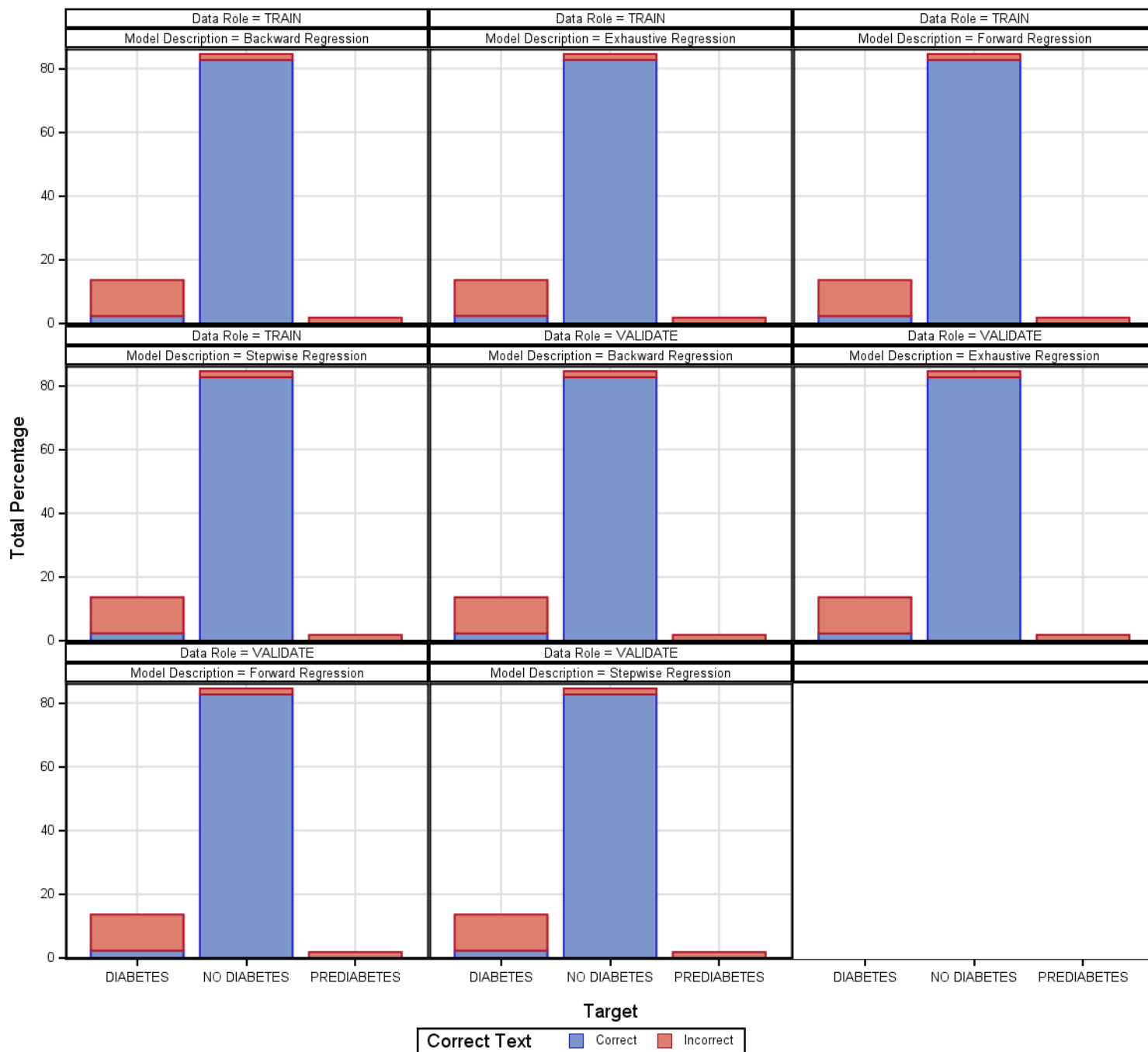
| Role | Level | Frequency Count | Name |
|--------|---------|-----------------|---------------------|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |

Node=MLR ModelComparison Fit Statistics Table

| Selected Model | Predecessor Node | Model Node | Model Description | Target Variable | Target Label | Selection Criterion: Valid: Misclassification Rate | Train: Average Squared Error | Train: Misclassification Rate | Train: Kolmogorov-Smirnov Statistic |
|----------------|------------------|------------|-----------------------|---------------------|--------------|--|------------------------------|-------------------------------|-------------------------------------|
| Y | Reg2 | Reg2 | Forward Regression | Diabetes_Prediction | | 0.15056 | 0.073718 | 0.15006 | 0.324 |
| | Reg3 | Reg3 | Backward Regression | Diabetes_Prediction | | 0.15056 | 0.073718 | 0.15006 | 0.324 |
| | Reg4 | Reg4 | Stepwise Regression | Diabetes_Prediction | | 0.15056 | 0.073718 | 0.15006 | 0.324 |
| | Reg | Reg | Exhaustive Regression | Diabetes_Prediction | | 0.15069 | 0.073600 | 0.14940 | 0.330 |

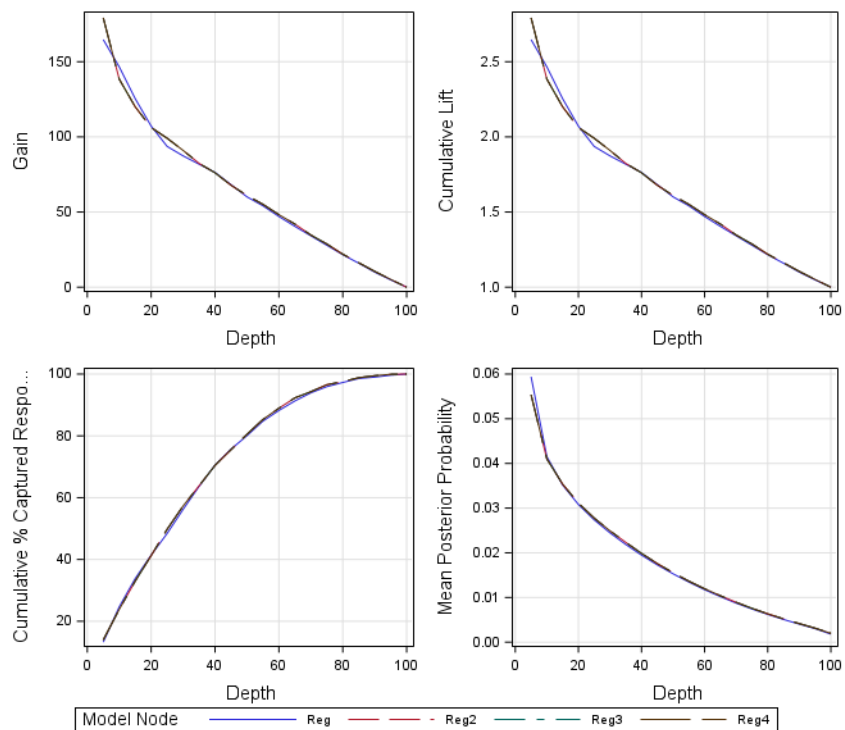
| Selected Model | Predecessor Node | Model Node | Model Description | Target Variable | Target Label | Selection Criterion: Valid: Misclassification Rate | Valid: Average Squared Error | Valid: Misclassification Rate | Valid: Kolmogorov-Smirnov Statistic |
|----------------|------------------|------------|-----------------------|---------------------|--------------|--|------------------------------|-------------------------------|-------------------------------------|
| Y | Reg2 | Reg2 | Forward Regression | Diabetes_Prediction | | 0.15056 | 0.073672 | 0.15056 | 0.307 |
| | Reg3 | Reg3 | Backward Regression | Diabetes_Prediction | | 0.15056 | 0.073672 | 0.15056 | 0.307 |
| | Reg4 | Reg4 | Stepwise Regression | Diabetes_Prediction | | 0.15056 | 0.073672 | 0.15056 | 0.307 |
| | Reg | Reg | Exhaustive Regression | Diabetes_Prediction | | 0.15069 | 0.073807 | 0.15069 | 0.304 |

SAS Enterprise Miner Report
Node=MLR ModelComparison
Classification Chart
TARGET='Diabetes_Prediction' and _TYPE_='PREDICTION'



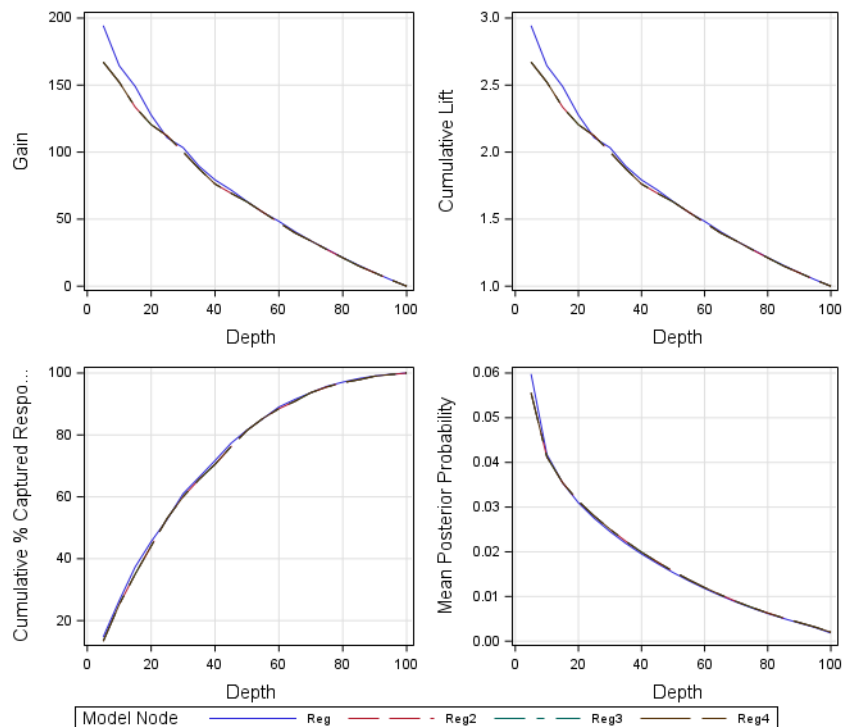
SAS Enterprise Miner Report

Node=MLR ModelComparison
Multiple Model Assessment Scores where DataRole=TEST
TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=MLR ModelComparison
Multiple Model Assessment Scores where DataRole=TRAIN
TARGET='Diabetes_Prediction'

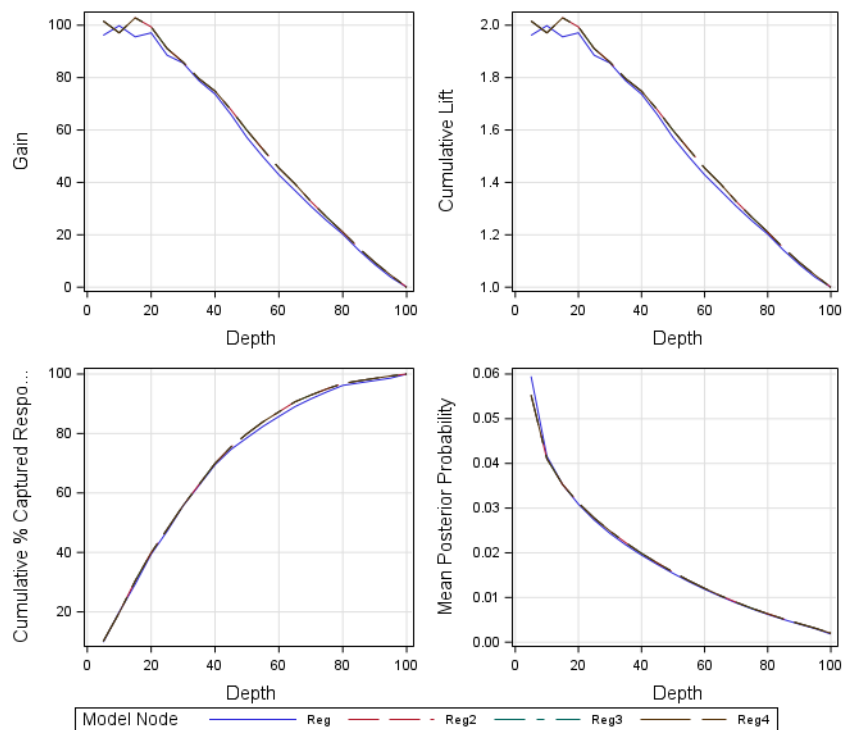


SAS Enterprise Miner Report

Node=MLR ModelComparison

Multiple Model Assessment Scores where DataRole=VALIDATE

TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree Model Comparison Summary

Node id = MdlComp
 Node label = RegTree Model Comparison
 Meta path = Ids3 => Part => Tree4 => MdlComp
 Notes =

Node=RegTree Model Comparison Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|-----------------------------|-------------------------------|---------|------------------------|---------|---------|----------------------|---------------------|---------|
| Component | ModelCompare | | NumberOfReportedLevels | 1E-6 | | SelectionData | DEFAULT | |
| AssessAllTargetLevels | N | | NumberOfBins | 20 | | SelectionDepth | 10 | |
| DecileBin | 20 | | ProfitEpsilon | 1E-6 | | SelectionTable | TRAIN | TABLE |
| HPCriteria | DEFAULT | | RecomputeAssess | N | | StatisticUsed | _VMISC_ | |
| LiftEpsilon | 1E-6 | | RocChart | Y | | TargetLabel | | |
| ModelCriteria | Valid: Misclassification Rate | | RocEpsilon | 0.01 | | TargetName | Diabetes_Prediction | |
| ModelDescription | RegTree B3D6 | | RoiEpsilon | 1E-6 | | classViyaCriteria | DEFAULT | |
| ModelId | Tree4 | | ScoreDistBin | 20 | | intervalViyaCriteria | DEFAULT | |
| NormalizeReportingVariables | Y | | SelectionCriteria | DEFAULT | | | | |

Node=RegTree Model Comparison Variable Summary

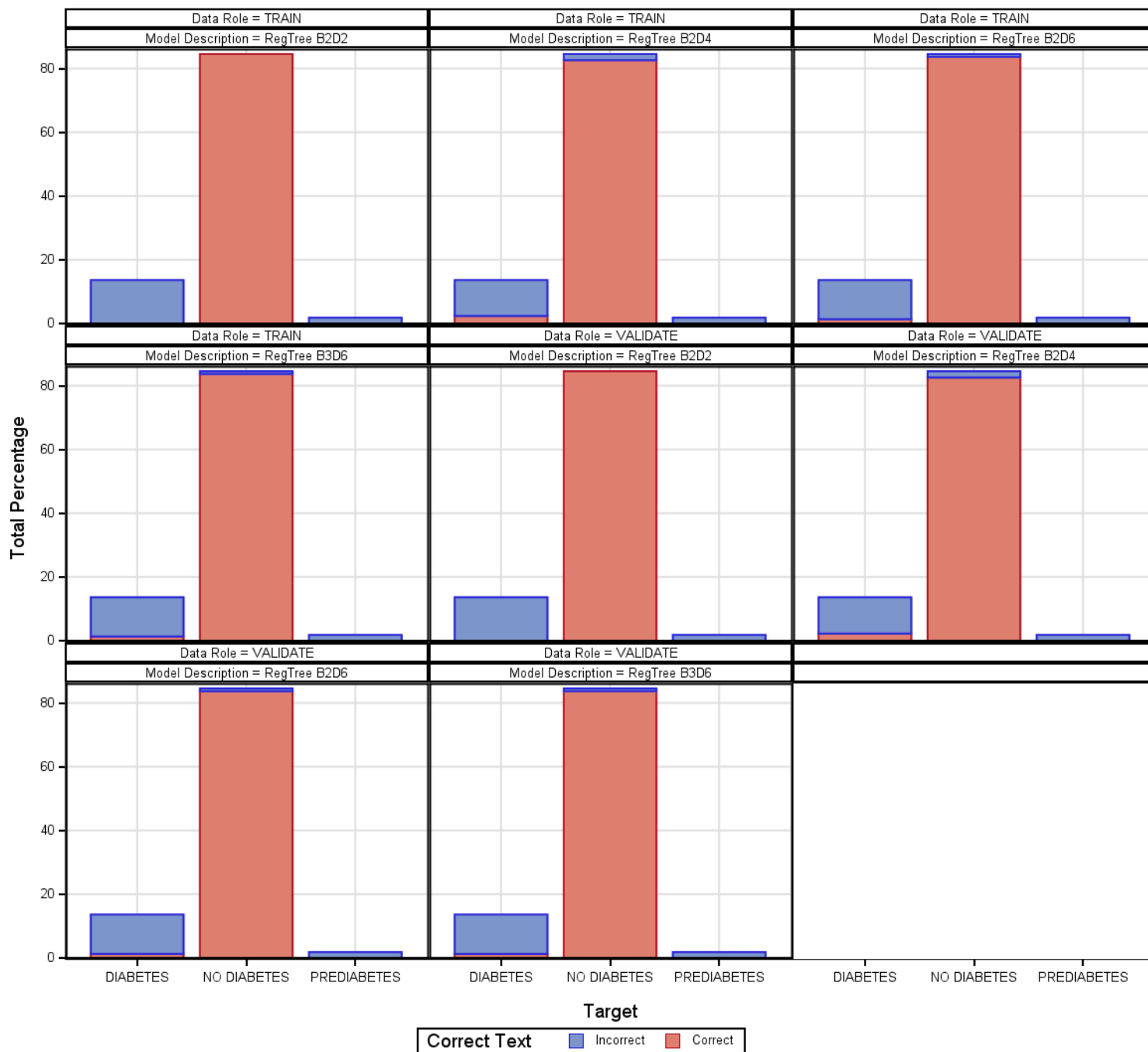
| Role | Level | Frequency Count | Name |
|--------|---------|-----------------|---------------------|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |

Node=RegTree Model Comparison Fit Statistics Table

| Selected Model | Predecessor Node | Model Node | Model Description | Target Variable | Target Label | Selection Criterion: Valid: Misclassification Rate | Train: Average Squared Error | Train: Misclassification Rate | Train: Kolmogorov-Smirnov Statistic |
|----------------|------------------|------------|-------------------|---------------------|--------------|--|------------------------------|-------------------------------|-------------------------------------|
| Y | Tree4 | Tree4 | RegTree B3D6 | Diabetes_Prediction | | 0.15053 | 0.079570 | 0.14959 | 0.212 |
| | Tree | Tree | RegTree B2D6 | Diabetes_Prediction | | 0.15061 | 0.079395 | 0.14979 | 0.207 |
| | Tree2 | Tree2 | RegTree B2D4 | Diabetes_Prediction | | 0.15201 | 0.079501 | 0.15041 | 0.207 |
| | Tree3 | Tree3 | RegTree B2D2 | Diabetes_Prediction | | 0.15463 | 0.088811 | 0.15464 | 0.000 |

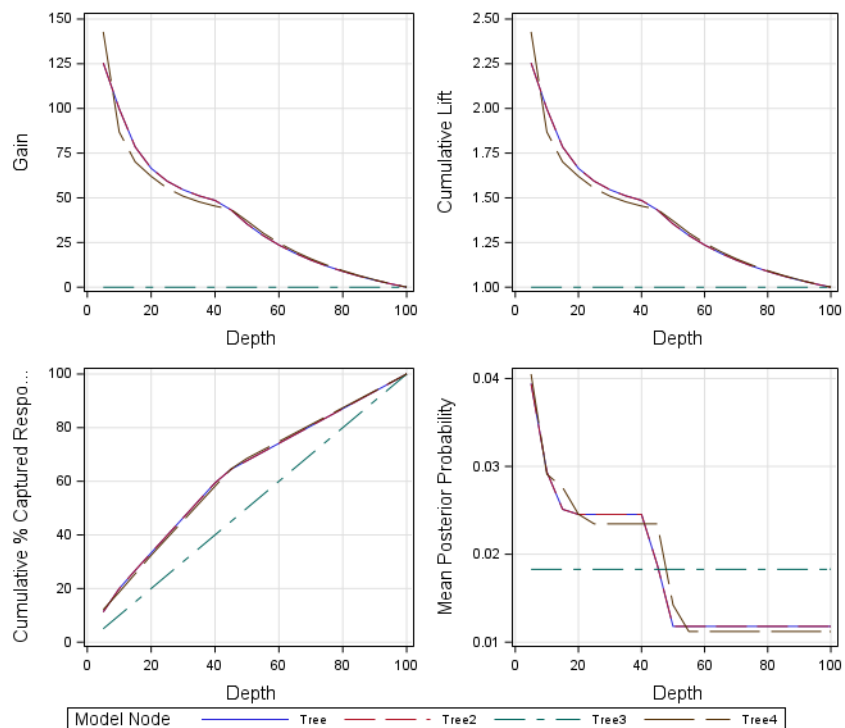
| Selected Model | Predecessor Node | Model Node | Model Description | Target Variable | Target Label | Selection Criterion: Valid: Misclassification Rate | Valid: Average Squared Error | Valid: Misclassification Rate | Valid: Kolmogorov-Smirnov Statistic |
|----------------|------------------|------------|-------------------|---------------------|--------------|--|------------------------------|-------------------------------|-------------------------------------|
| Y | Tree4 | Tree4 | RegTree B3D6 | Diabetes_Prediction | | 0.15053 | 0.079877 | 0.15053 | 0.204 |
| | Tree | Tree | RegTree B2D6 | Diabetes_Prediction | | 0.15061 | 0.079762 | 0.15061 | 0.210 |
| | Tree2 | Tree2 | RegTree B2D4 | Diabetes_Prediction | | 0.15201 | 0.079872 | 0.15201 | 0.210 |
| | Tree3 | Tree3 | RegTree B2D2 | Diabetes_Prediction | | 0.15463 | 0.088810 | 0.15463 | 0.000 |

SAS Enterprise Miner Report
Node=RegTree Model Comparison
Classification Chart
TARGET='Diabetes_Prediction' and _TYPE_='PREDICTION'



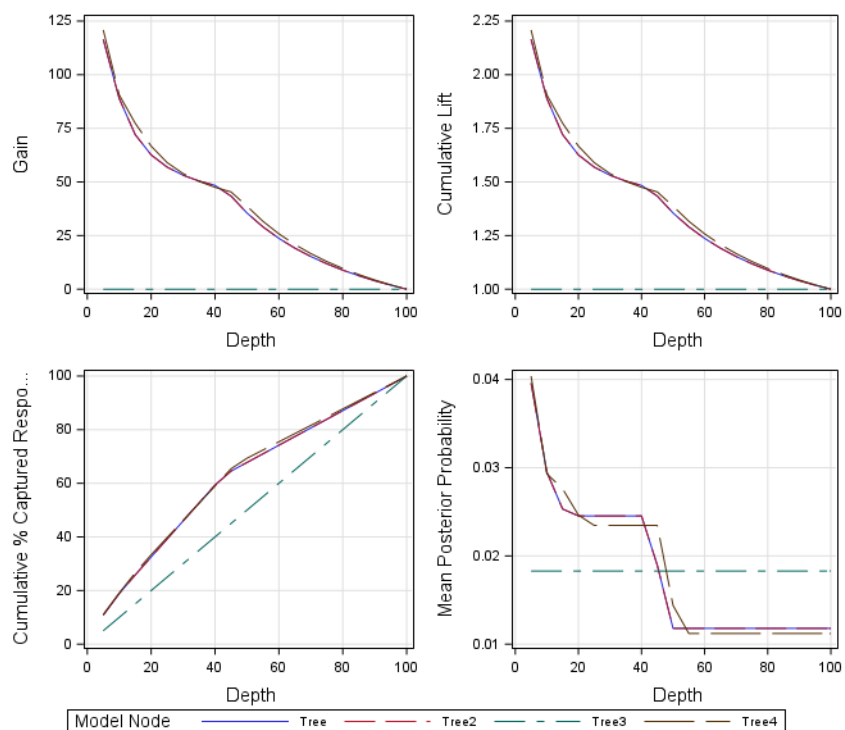
SAS Enterprise Miner Report

Node=RegTree Model Comparison
Multiple Model Assessment Scores where DataRole=TEST
TARGET='Diabetes_Prediction'



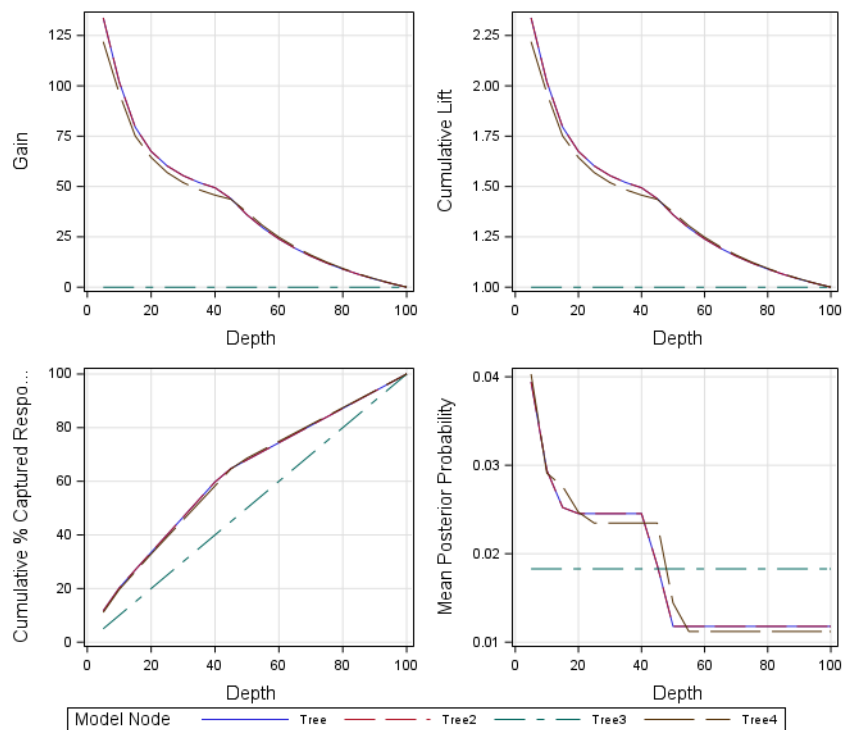
SAS Enterprise Miner Report

Node=RegTree Model Comparison
Multiple Model Assessment Scores where DataRole=TRAIN
TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=RegTree Model Comparison
Multiple Model Assessment Scores where DataRole=VALIDATE
TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=SCOREDIABETES_TRAIN
Summary

Node id = Ids4
Node label = SCOREDIABETES_TRAIN
Meta path = Ids4
Notes =

Node=SCOREDIABETES_TRAIN
Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|------------------------------|------------------------------|---------|----------------------|--------------------|---------|-------------------|---------------------|---------|
| Component | DataSource | | DsCreatedBy | shre2 | | NBytes | 9307136 | . |
| ApplyIntervalLevelLowerLimit | Y | | DsId | scorediabetestrain | | NCols | 22 | . |
| ApplyMaxClassLevels | Y | | DsModifiedBy | shre2 | | NObs | 50144 | . |
| ApplyMaxPercentMissing | Y | | DsModifyDate | 2025130005.4 | | NewTable | | |
| CMeta | WORK.M0_AXJ7R | | DsSampleName | | | NewVariableRole | REJECT | |
| ComputeStatistics | N | | DsSampleSize | | | OutputType | VIEW | |
| DBPassThrough | Y | | DsSampleSizeType | | | Role | RAW | TRAIN |
| Data | ZETADATA.SCOREDIABETES_TRAIN | | DsScope | LOCAL | | Sample | D | |
| DataSelection | DATASOURCE | | IdentifyEmptyColumns | Y | | SampleSizeObs | 10000 | |
| DataSource | scorediabetestrain | | IntervalLowerLimit | 20 | | SampleSizePercent | 20 | |
| DataSourceRole | RAW | | Library | ZETADATA | | SampleSizeType | PERCENT | |
| Description | | | MaxClassLevels | 20 | | Scope | LOCAL | |
| DropMapVariables | Y | | MaxPercentMissing | 50 | | Segment | | |
| DsCreateDate | 2025130005.1 | | MetaAdvisor | BASIC | | Table | SCOREDIABETES_TRAIN | |

Node=SCOREDIABETES_TRAIN
Data Attributes

| Attribute | Value | Attribute | Value | Attribute | Value |
|------------|---------------------|----------------|--------------------|--------------|----------|
| Data Name | SCOREDIABETES_TRAIN | Date Created | 04Mar2024:00:05:17 | Data Size | 9307136 |
| Data Type | DATA | Date Modified | 04Mar2024:00:05:17 | Role | RAW |
| Data Label | | Number Rows | 50144 | Segment | |
| Engine | BASE | Number Columns | 22 | Data Library | ZETADATA |

Node=SCOREDIABETES_TRAIN
Variables List

| Name | Label | Role | Level | Type | Length | Format | Creator |
|----------------------|----------------------|--------|----------|------|--------|--------|---------|
| Age | Age | INPUT | INTERVAL | N | 8 | BEST. | |
| AnyHealthcare | AnyHealthcare | INPUT | BINARY | N | 8 | BEST. | |
| BMI | BMI | INPUT | INTERVAL | N | 8 | BEST. | |
| CholCheck | CholCheck | INPUT | BINARY | N | 8 | BEST. | |
| Diabetes_Prediction | | TARGET | NOMINAL | C | 11 | | |
| DiffWalk | DiffWalk | INPUT | BINARY | N | 8 | BEST. | |
| Education | Education | INPUT | ORDINAL | N | 8 | BEST. | |
| Fruits | Fruits | INPUT | BINARY | N | 8 | BEST. | |
| GenHlth | GenHlth | INPUT | ORDINAL | N | 8 | BEST. | |
| HeartDiseaseorAttack | HeartDiseaseorAttack | INPUT | BINARY | N | 8 | BEST. | |
| HighBP | HighBP | INPUT | BINARY | N | 8 | BEST. | |

| Name | Label | Role | Level | Type | Length | Format | Creator |
|-------------------|-------------------|-------|---------|------|--------|--------|---------|
| HighChol | HighChol | INPUT | BINARY | N | 8 | BEST. | |
| HvyAlcoholConsump | HvyAlcoholConsump | INPUT | BINARY | N | 8 | BEST. | |
| Income | Income | INPUT | ORDINAL | N | 8 | BEST. | |
| MentHlth | MentHlth | INPUT | ORDINAL | N | 8 | BEST. | |
| NoDocbcCost | NoDocbcCost | INPUT | BINARY | N | 8 | BEST. | |
| PhysActivity | PhysActivity | INPUT | BINARY | N | 8 | BEST. | |
| PhysHlth | PhysHlth | INPUT | ORDINAL | N | 8 | BEST. | |
| Sex | Sex | INPUT | BINARY | N | 8 | BEST. | |
| Smoker | Smoker | INPUT | BINARY | N | 8 | BEST. | |
| Stroke | Stroke | INPUT | BINARY | N | 8 | BEST. | |
| Veggies | Veggies | INPUT | BINARY | N | 8 | BEST. | |

SAS Enterprise Miner Report

Node=Predict Model Comparison
Summary

Node id = MdlComp3
Node label = Predict Model Comparison
Meta path = Ids3 => Part => Tree4 => MdlComp => MdlComp3
Notes =

Node=Predict Model Comparison
Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|-----------------------------|-------------------------------|---------|------------------------|---------|---------|----------------------|---------------------|---------|
| Component | ModelCompare | | NumberOfReportedLevels | 1E-6 | | SelectionData | DEFAULT | |
| AssessAllTargetLevels | N | | NumberOfBins | 20 | | SelectionDepth | 10 | |
| DecileBin | 20 | | ProfitEpsilon | 1E-6 | | SelectionTable | TRAIN | TABLE |
| HPCriteria | DEFAULT | | RecomputeAssess | N | | StatisticUsed | _VMISC_ | |
| LiftEpsilon | 1E-6 | | RocChart | Y | | TargetLabel | | |
| ModelCriteria | Valid: Misclassification Rate | | RocEpsilon | 0.01 | | TargetName | Diabetes_Prediction | |
| ModelDescription | RegTree B3D6 | | RoiEpsilon | 1E-6 | | classViyaCriteria | DEFAULT | |
| ModelId | Tree4 | | ScoreDistBin | 20 | | intervalViyaCriteria | DEFAULT | |
| NormalizeReportingVariables | Y | | SelectionCriteria | DEFAULT | | | | |

Node=Predict Model Comparison
Variable Summary

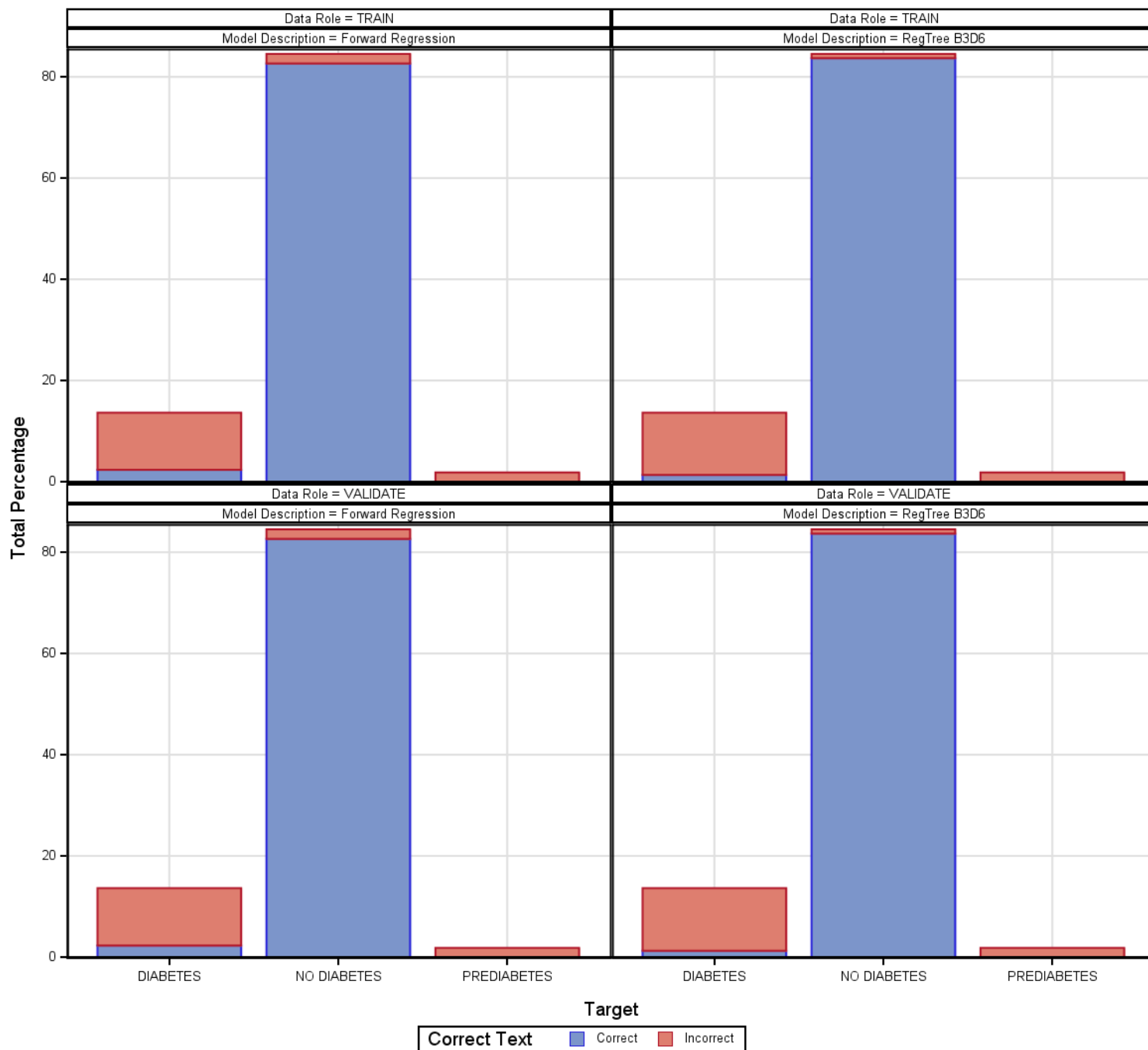
| Role | Level | Frequency Count | Name |
|--------|---------|-----------------|---------------------|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |

Node=Predict Model Comparison
Fit Statistics Table

| Selected Model | Predecessor Node | Model Node | Model Description | Target Variable | Target Label | Selection Criterion: Valid: Misclassification Rate | Train: Average Squared Error | Train: Misclassification Rate | Train: Kolmogorov-Smirnov Statistic |
|----------------|------------------|------------|--------------------|---------------------|--------------|--|------------------------------|-------------------------------|-------------------------------------|
| Y | MdlComp | Tree4 | RegTree B3D6 | Diabetes_Prediction | | 0.15053 | 0.079570 | 0.14959 | 0.212 |
| | MdlComp2 | Reg2 | Forward Regression | Diabetes_Prediction | | 0.15056 | 0.073718 | 0.15006 | 0.324 |

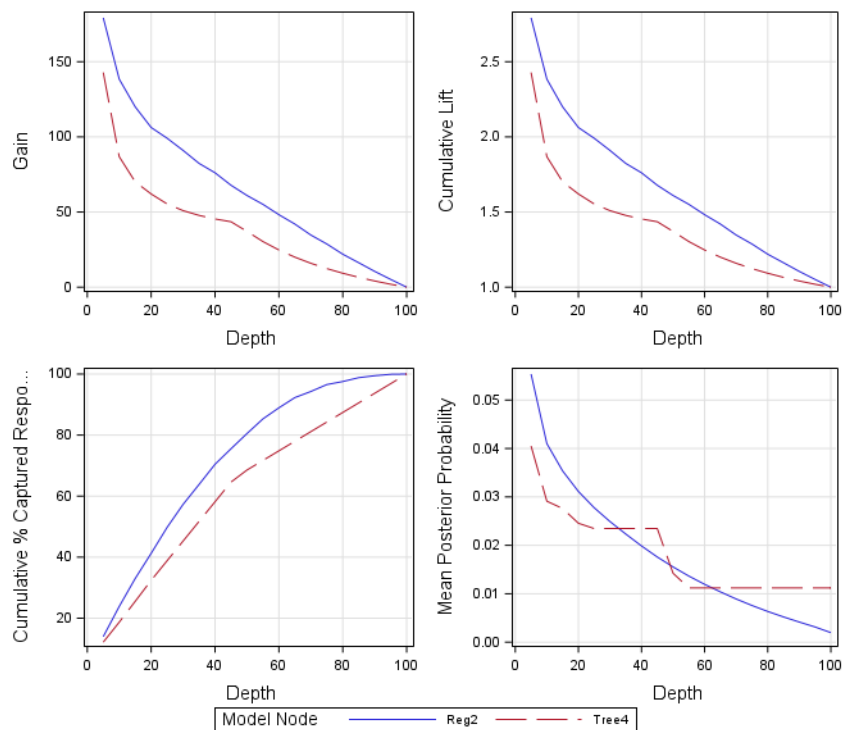
| Selected Model | Predecessor Node | Model Node | Model Description | Target Variable | Target Label | Selection Criterion: Valid: Misclassification Rate | Valid: Average Squared Error | Valid: Misclassification Rate | Valid: Kolmogorov-Smirnov Statistic |
|----------------|------------------|------------|--------------------|---------------------|--------------|--|------------------------------|-------------------------------|-------------------------------------|
| Y | MdlComp | Tree4 | RegTree B3D6 | Diabetes_Prediction | | 0.15053 | 0.079877 | 0.15053 | 0.204 |
| | MdlComp2 | Reg2 | Forward Regression | Diabetes_Prediction | | 0.15056 | 0.073672 | 0.15056 | 0.307 |

SAS Enterprise Miner Report
Node=Predict Model Comparison
Classification Chart
TARGET='Diabetes_Prediction' and _TYPE_='PREDICTION'



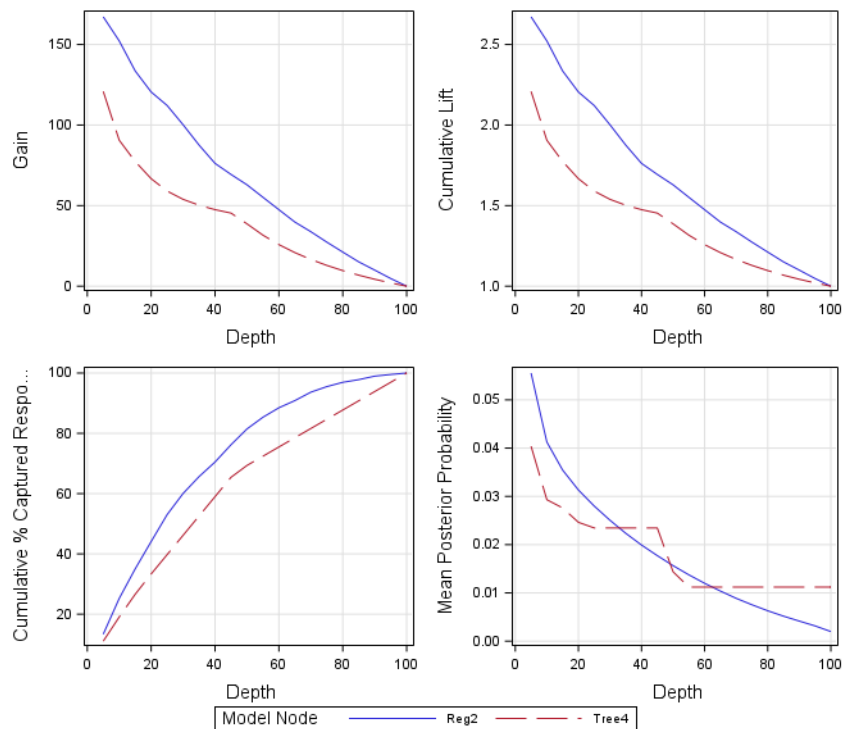
SAS Enterprise Miner Report

Node=Predict Model Comparison
Multiple Model Assessment Scores where DataRole=TEST
TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=Predict Model Comparison
Multiple Model Assessment Scores where DataRole=TRAIN
TARGET='Diabetes_Prediction'

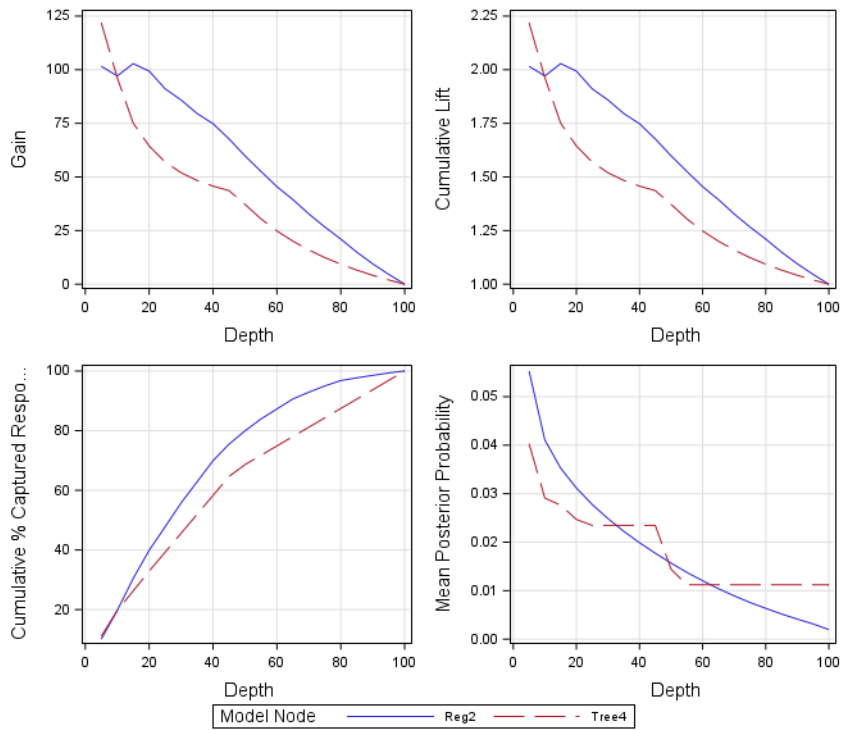


SAS Enterprise Miner Report

Node=Predict Model Comparison

Multiple Model Assessment Scores where DataRole=VALIDATE

TARGET='Diabetes_Prediction'



SAS Enterprise Miner Report

Node=Score Summary

Node id = Score
Node label = Score
Meta path = Ids3 => Part => Tree4 => MdlComp => MdlComp3 => Score
Notes =

Node=Score Properties

| Property | Value | Default | Property | Value | Default | Property | Value | Default |
|--------------------|-------|---------|---------------|-------|---------|----------------|---------|---------|
| Component | Score | | HideInput | Y | | JScore | N | |
| CScore | N | | HideOther | Y | | OptimizedCode | Y | |
| FixedOutputNames | Y | | HidePredict | Y | | OutputType | VIEW | |
| GraphReports | Y | | HideRejected | Y | | PackageName | DEFAULT | |
| HideAssess | Y | | HideResidual | Y | | PreferenceName | | |
| HideClassification | Y | | HideTarget | Y | | ScoreTest | N | |
| HideFreq | Y | | HideVariables | N | | ScoreValidate | N | |

Node=Score Variable Summary

| Role | Level | Frequency Count | Name |
|---------|---------|--------------------|---------------------|
| TARGET | NOMINAL | 1 | Diabetes_Prediction |
| SEGMENT | NOMINAL | 1 | _NODE_ |

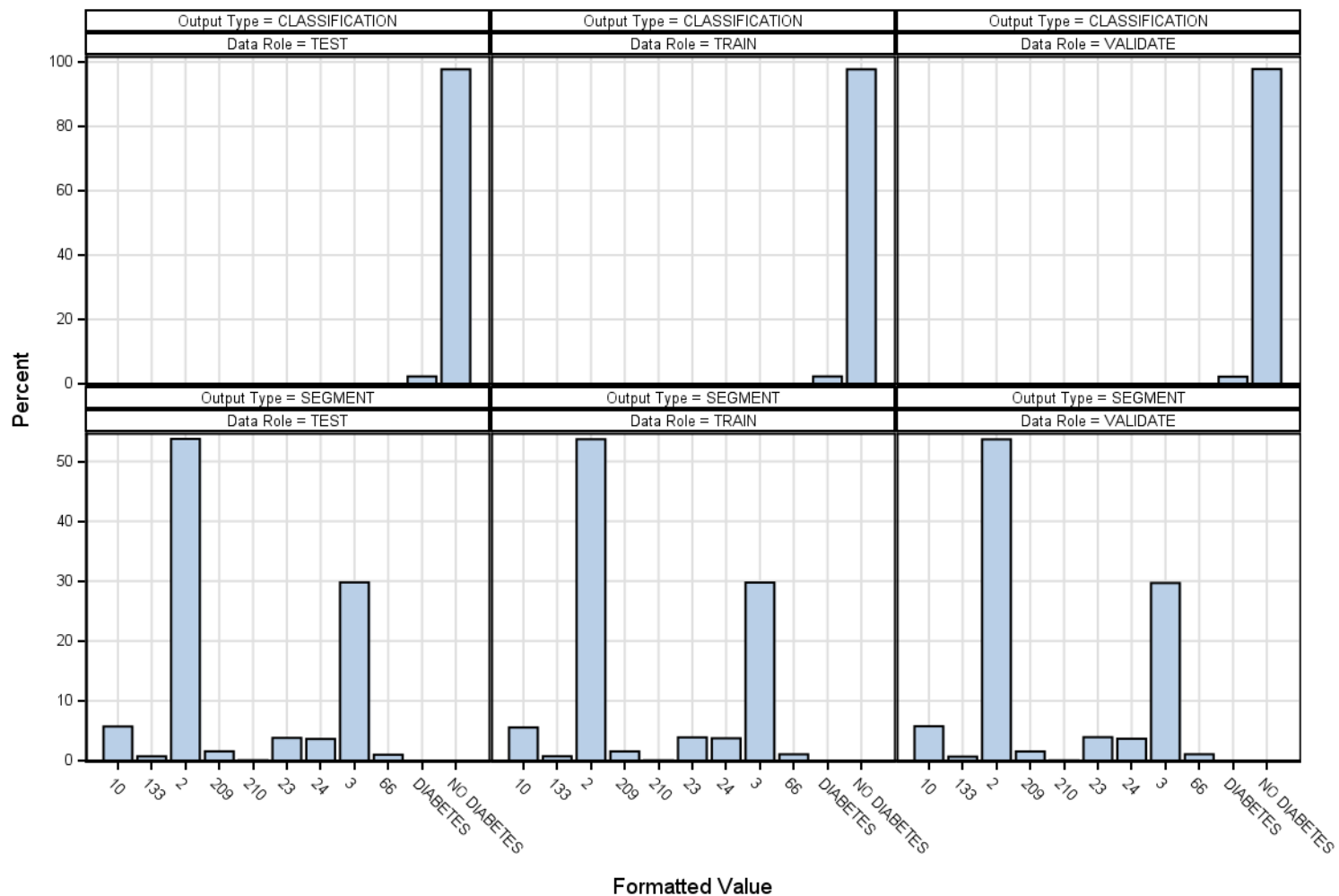
Node=Score Output Variables

| Variable Name | Creator | Variable Label | Function | Type |
|----------------------------------|---------|--|----------------|------|
| EM_CLASSIFICATION | Score | Prediction for Diabetes_Prediction | CLASSIFICATION | C |
| EM_EVENTPROBABILITY | Score | Probability for level PREDIABETES of Diabetes_Prediction | PREDICT | N |
| EM_PROBABILITY | Score | Probability of Classification | PREDICT | N |
| EM_SEGMENT | Score | Node | TRANSFORM | N |
| I_Diabetes_Prediction | Tree4 | Into: Diabetes_Prediction | CLASSIFICATION | C |
| P_Diabetes_Predictiondiabetes | Tree4 | Predicted: Diabetes_Prediction=diabetes | PREDICT | N |
| P_Diabetes_Predictionno_diabetes | Tree4 | Predicted: Diabetes_Prediction=no diabetes | PREDICT | N |
| P_Diabetes_Predictionprediabetes | Tree4 | Predicted: Diabetes_Prediction=prediabetes | PREDICT | N |
| Q_Diabetes_Predictiondiabetes | Tree4 | Unadjusted P: Diabetes_Prediction=diabetes | TRANSFORM | N |
| Q_Diabetes_Predictionno_diabetes | Tree4 | Unadjusted P: Diabetes_Prediction=no diabetes | TRANSFORM | N |
| Q_Diabetes_Predictionprediabetes | Tree4 | Unadjusted P: Diabetes_Prediction=prediabetes | TRANSFORM | N |
| U_Diabetes_Prediction | Tree4 | Unnormalized Into: Diabetes_Prediction | CLASSIFICATION | C |
| V_Diabetes_Predictiondiabetes | Tree4 | Validated: Diabetes_Prediction=diabetes | PREDICT | N |
| V_Diabetes_Predictionno_diabetes | Tree4 | Validated: Diabetes_Prediction=no diabetes | PREDICT | N |
| V_Diabetes_Predictionprediabetes | Tree4 | Validated: Diabetes_Prediction=prediabetes | PREDICT | N |
| _NODE_ | Tree4 | Node | TRANSFORM | N |
| _WARN_ | Tree4 | Warnings | ASSESS | C |

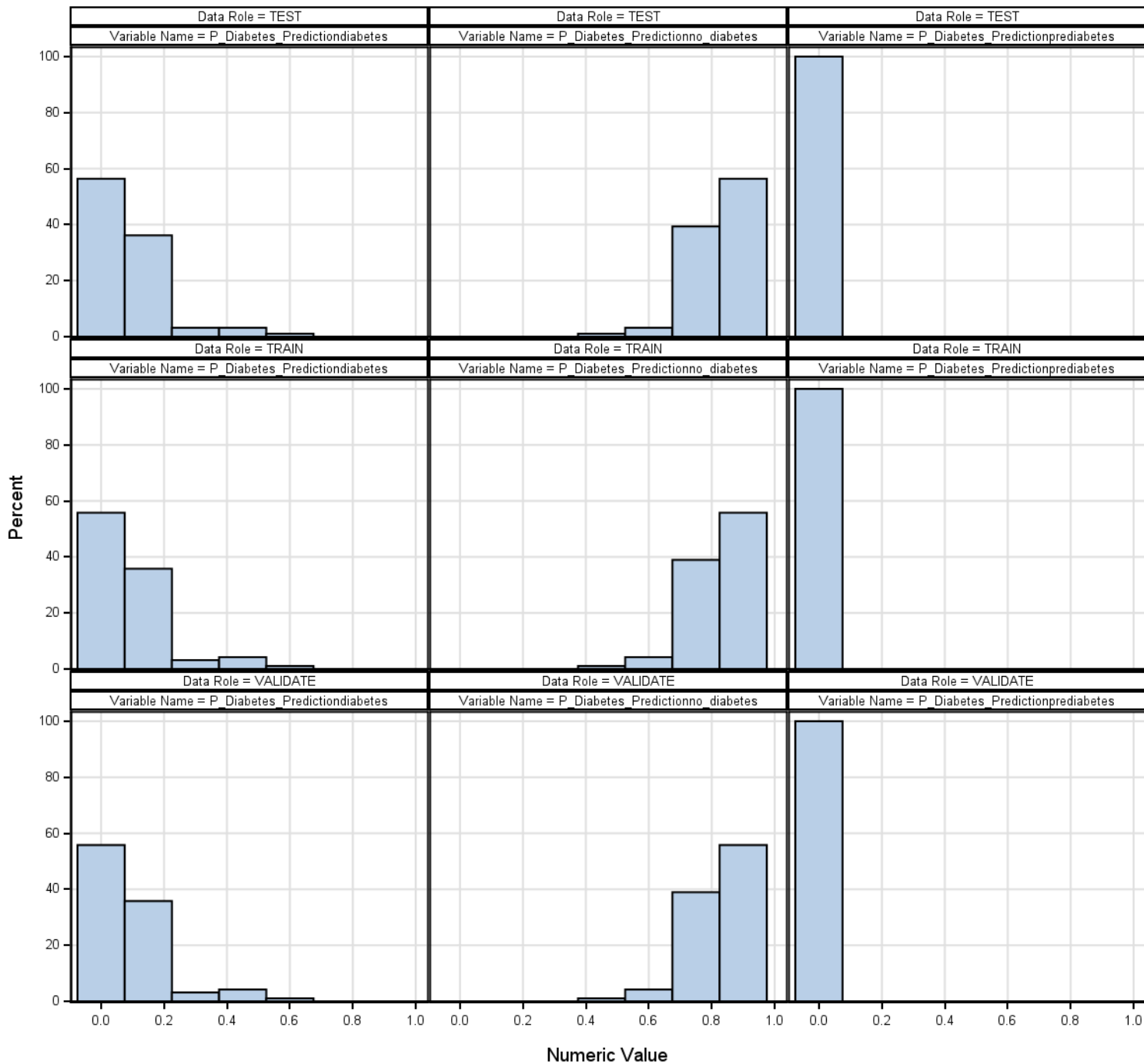
SAS Enterprise Miner Report

Node=Score

Bar Chart



SAS Enterprise Miner Report
Node=Score
Histogram



End of Report