J48 pruned tree

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size\_uniformity = S

| bare\_nucleoli = S

| | normal\_nucleoli = S: Benign (416.0/3.0)

| | normal\_nucleoli = M

| | | bland\_chromatin = S: Benign (3.0)

| | | bland\_chromatin = H: Benign (0.0)

| | | bland\_chromatin = M: Malignant (2.0)

| | normal\_nucleoli = H

| | | shape\_uniformity = S: Benign (2.0)

| | | shape\_uniformity = H: Benign (0.0)

| | | shape\_uniformity = M: Malignant (2.0)

| bare\_nucleoli = M

| | clump\_thickness = M

| | | epithelial\_size = S: Malignant (6.0/1.0)

| | | epithelial\_size = M: Benign (3.0/1.0)

| | | epithelial\_size = H: Malignant (0.0)

| | clump\_thickness = S: Benign (11.0/1.0)

| | clump\_thickness = H: Malignant (4.0)

| bare\_nucleoli = H: Malignant (21.0/2.0)

size\_uniformity = H: Malignant (101.0/2.0)

size\_uniformity = M: Malignant (111.0/8.0)