

Initial Results for Meeting.

— using RandomForest.

Before Balancing:

- *Baseline Function Accuracy:*
 - 0.4603775 — `sum(dog_total$Outcome_Type == Adoption)/nrows(dog_total)`
- *Random Forest Accuracy:*
 - 0.9413287

Call:

```
randomForest(formula = Outcome.Type ~ ., data = train)
```

Type of random forest: classification

Number of trees: 500

No. of variables tried at each split: 3

OOB estimate of error rate: 5.36%

Confusion matrix:

	Adoption	Disposal	Euthanasia	Missing	Return to Owner	Rto-Adopt	Transfer
Adoption	15775	0	0	0	384	0	0
Disposal	5	1	0	0	4	0	0
Euthanasia	2	0	1071	0	2	0	0
Missing	1	0	0	0	3	0	4
Return to Owner	1330	0	1	0	8897	0	0
Rto-Adopt	132	0	0	0	15	0	0
Transfer	0	0	0	0	0	0	7530

class.error

Adoption	0.02376385
Disposal	0.90000000
Euthanasia	0.00372093
Missing	1.00000000
Return to Owner	0.13013297
Rto-Adopt	1.00000000
Transfer	0.00000000

Purebred.or.Mixed	Gender	Intake.Condition	Found.Location	Gen.Color
36.57362	61.81315	216.83973	220.43282	342.08906
AKC.Group	Outcome.Month	Age.upon.Outcome	Sex.upon.Outcome	Sex.Difference
411.89071	524.43941	999.82782	1002.50222	1450.98899
Intake.Type	Days.to.Outcome	Outcome.Subtype		
2704.68630	4393.47602	9656.09670		

^^ MeanDecreaseGini

After Balancing:

- *Random Forest Accuracy:*
 - 0.9386536

Distribution of Classification Classes:

Class	Total Entries
Missing	16
Disposal	15
RTO	14571
RTO-Adopt	216
Transfer	10747
Euthanasia	1537
Adopt	23122

Call:

```
randomForest(formula = Outcome.Type ~ ., data = train_smote)
  Type of random forest: classification
    Number of trees: 500
```

No. of variables tried at each split: 3

OOB estimate of error rate: 3.11%

Confusion matrix:

	Adoption	Disposal	Euthanasia	Missing	Return to Owner	Rto-Adopt	Transfer
Adoption	7517	23	0	0	134	0	0
Disposal	7	8289	0	0	20	0	0
Euthanasia	3	1	497	0	1	0	0
Missing	0	0	0	2	2	0	1
Return to Owner	486	26	1	0	4253	0	0
Rto-Adopt	63	0	0	0	7	11	0
Transfer	0	0	0	0	0	0	3586

	class.error
Adoption	0.020458692
Disposal	0.003246753
Euthanasia	0.009960159
Missing	0.600000000
Return to Owner	0.107637432
Rto-Adopt	0.864197531
Transfer	0.000000000

```
> tree_test[[1]]$importance[order(tree_test[[1]]$importance),]
```

Purebred.or.Mixed	Gender	Found.Location	Gen.Color	AKC.Group
47.03807	138.40770	414.12539	506.20640	621.11206
Intake.Condition	Intake.Type	Age.upon.Outcome	Sex.Difference	Outcome.Month
1051.52949	1245.95766	1296.16407	1421.49660	1445.26952
Days.to.Outcome	Sex.upon.Outcome	Outcome.Subtype		
2461.48005	2487.01568	4716.51258		