Gene expression analysis in Puleo seq

Ryan Kawalerski

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Contents
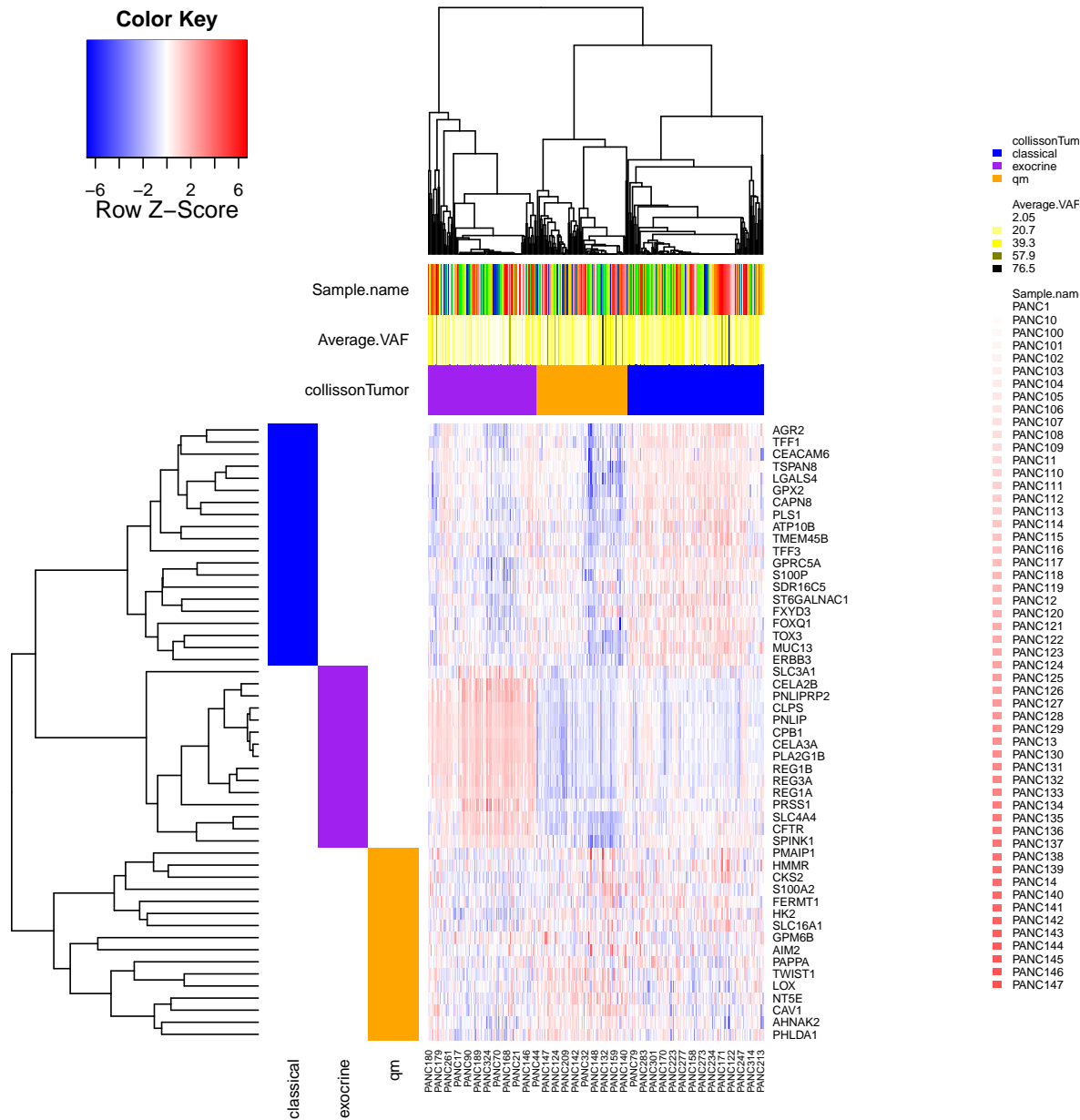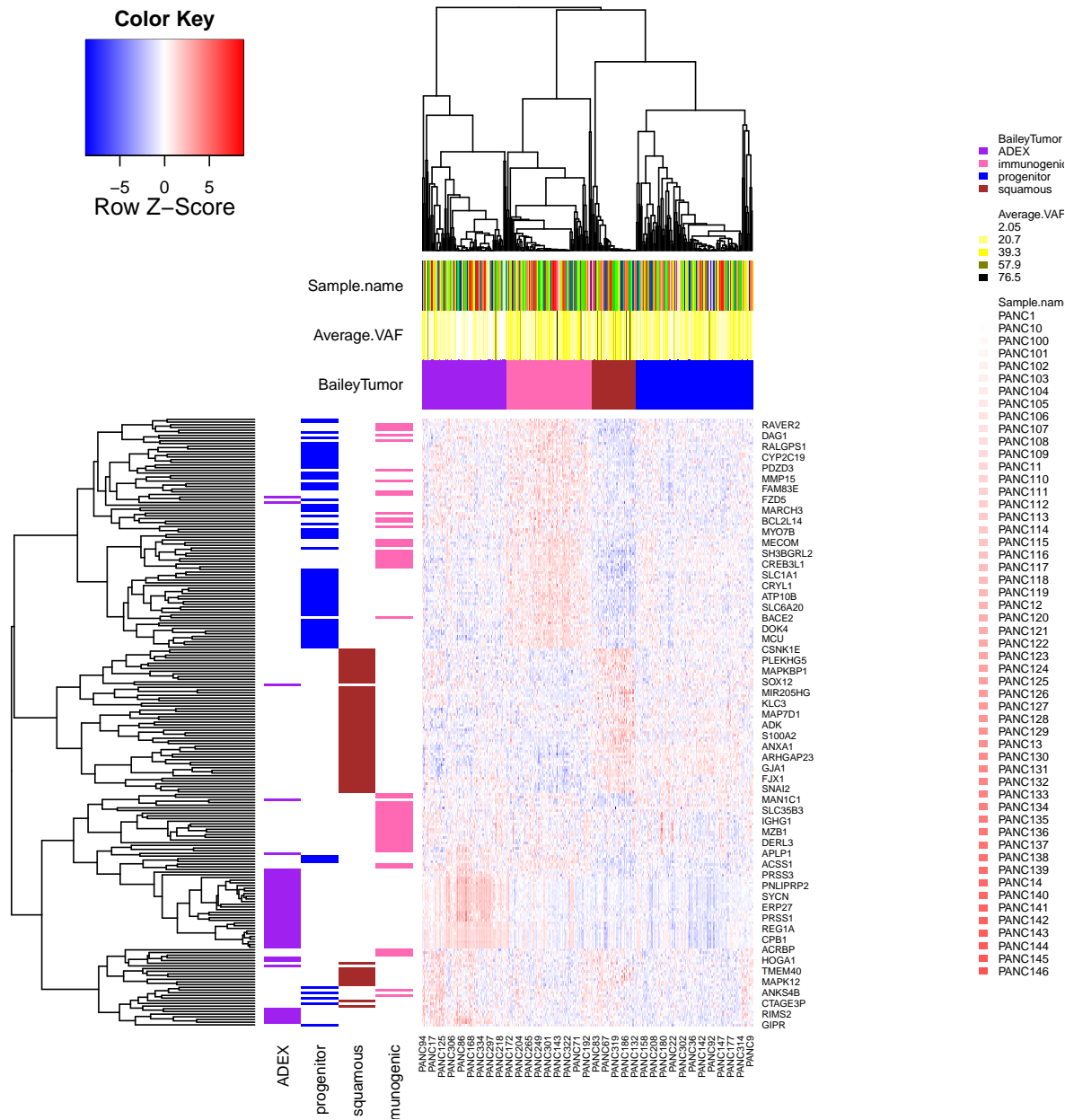
Assign dataset and remove nonvariant genes

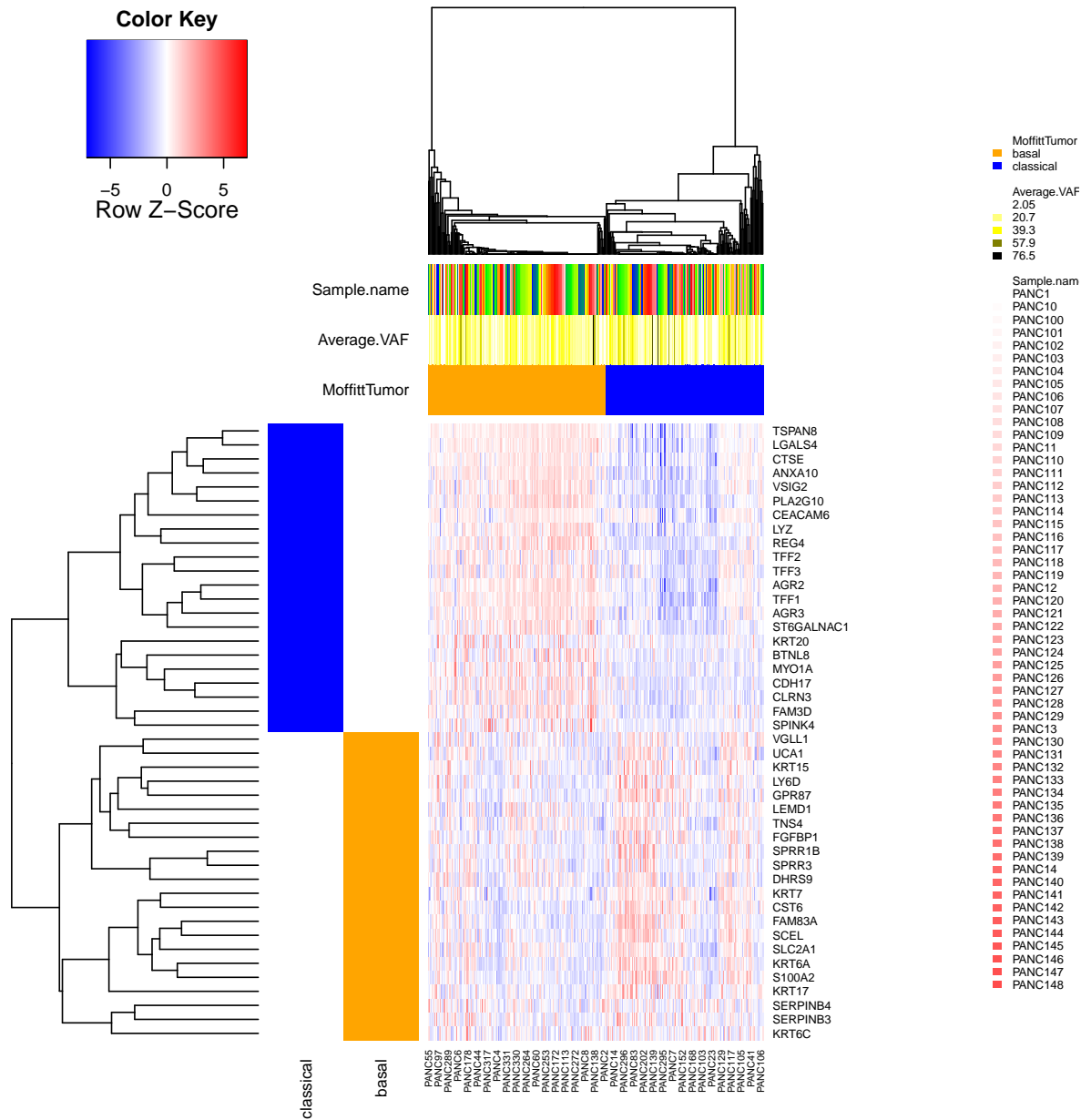## Add genesets to dataset sample info

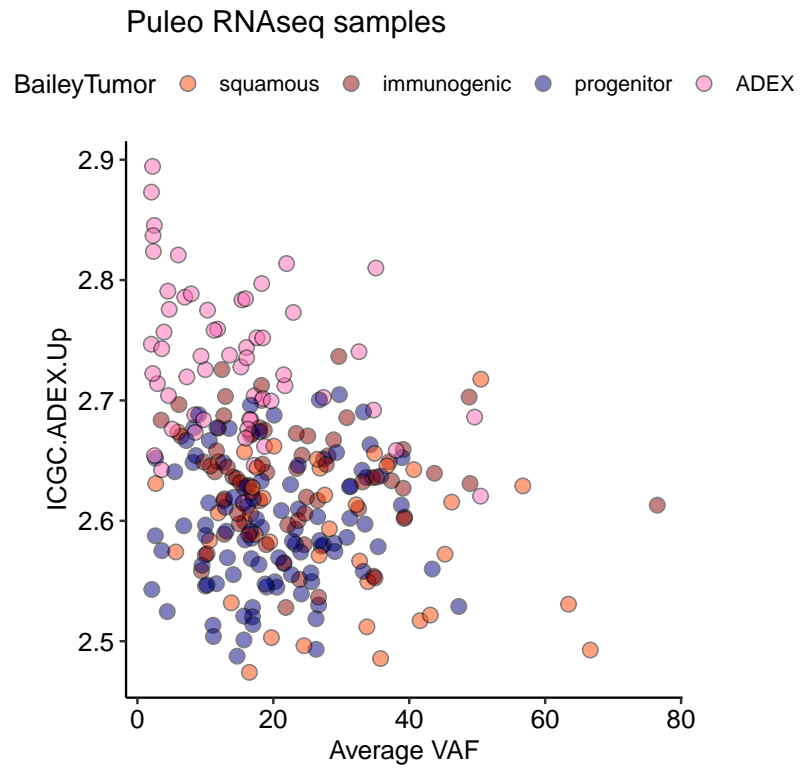## Consensus clustering

Collisson subtypes

Bailey subtypes

Moffitt subtypes

Gene signature expression by purity and subtypes
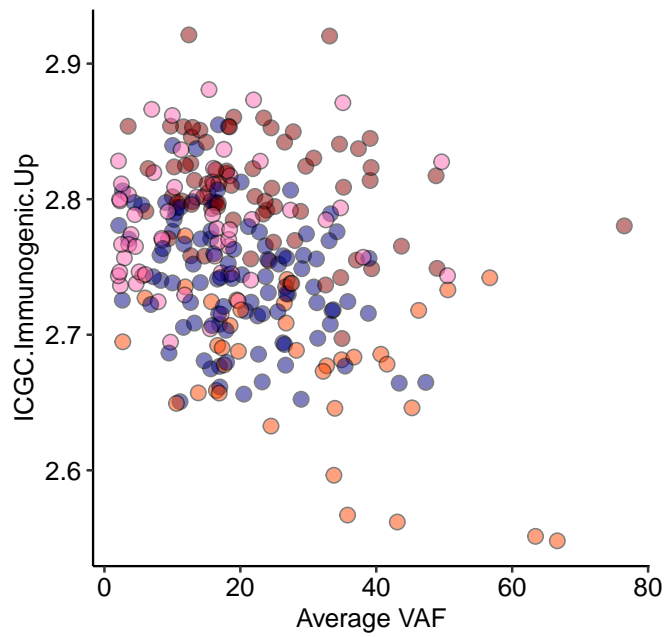
Signatures and average VAF colored by Bailey subtypes

Puleo RNAseq samples


```
## [1] -0.27207
## [1] 6.22e-06
```
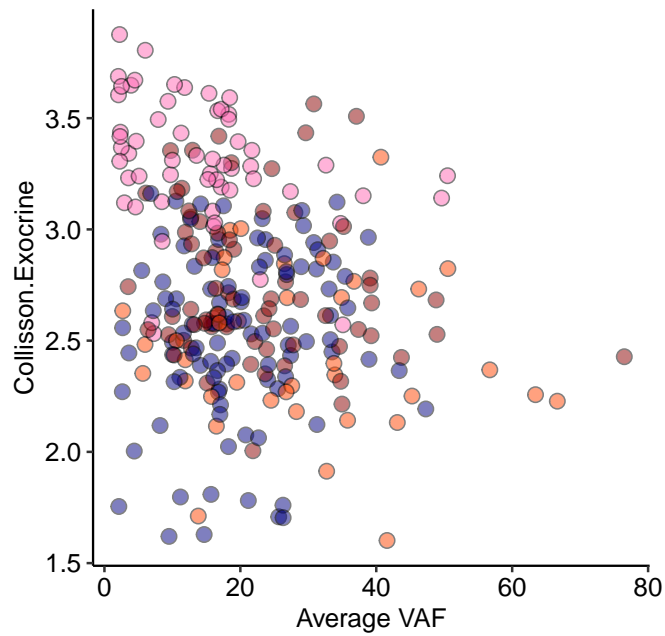
Puleo RNAseq samples

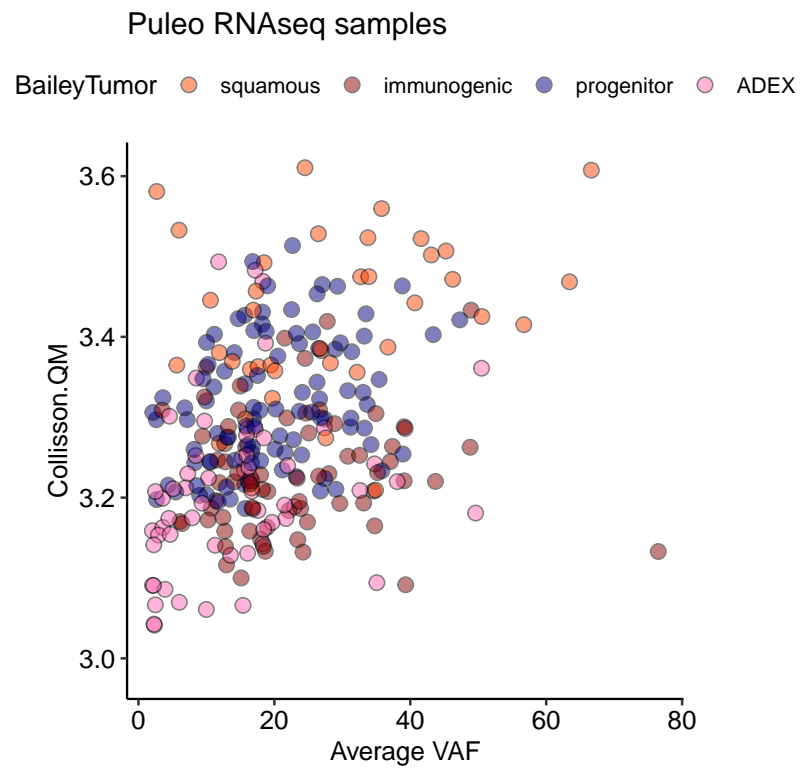BaileyTumor ⬤ squamous ⬤ immunogenic ⬤ progenitor ⬤ ADEX


```
## [1] -0.27956
## [1] 3.35e-06
```

Puleo RNAseq samples
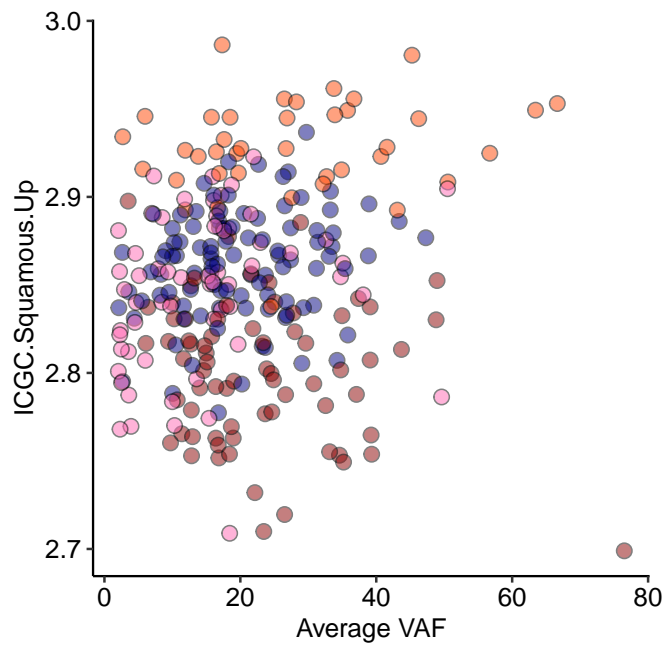
BaileyTumor ⬤ squamous ⬤ immunogenic ⬤ progenitor ⬤ ADEX

```
## [1] -0.23084
## [1] 0.00013729
```


Puleo RNAseq samples

```
## [1] 0.32705
## [1] 4e-08
```

Puleo RNAseq samples

[1] 0.13103
[1] 0.03201589

Puleo RNAseq samples

```
## [1] 0.28846
## [1] 1.56e-06
```

Puleo RNAseq samples


```
## [1] -0.22488
## [1] 0.00020575
```
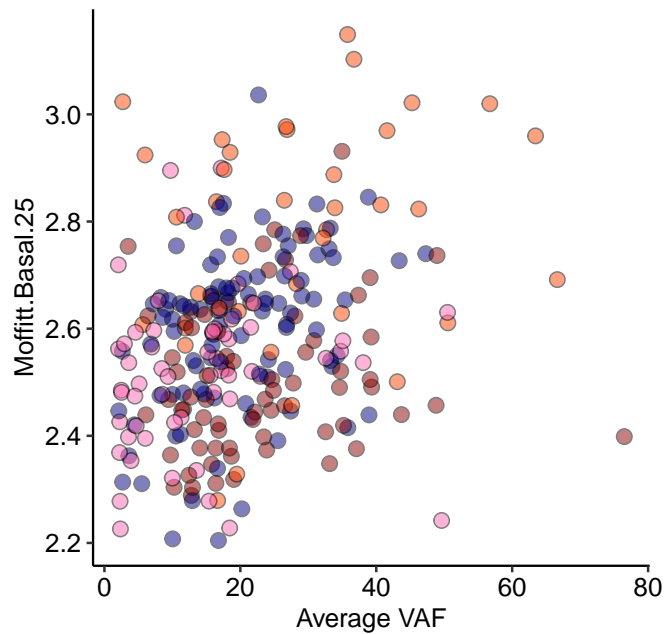
Puleo RNAseq samples

BaileyTumor ● squamous ● immunogenic ● progenitor ● ADEX


```
## [1] 0.24632
## [1] 4.571e-05
```

Puleo RNAseq samples

BaileyTumor ● squamous ● immunogenic ● progenitor ● ADEX
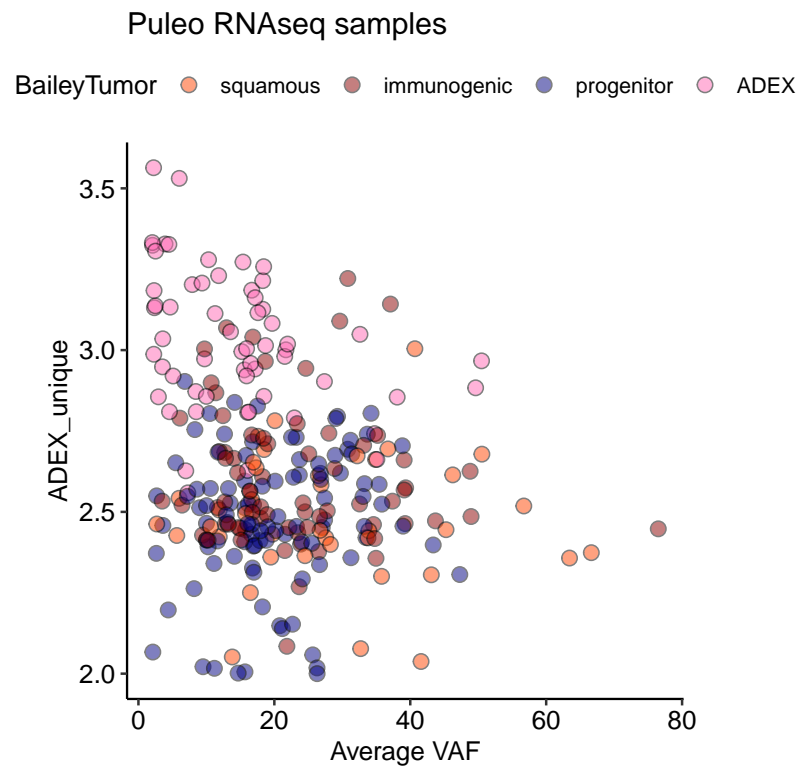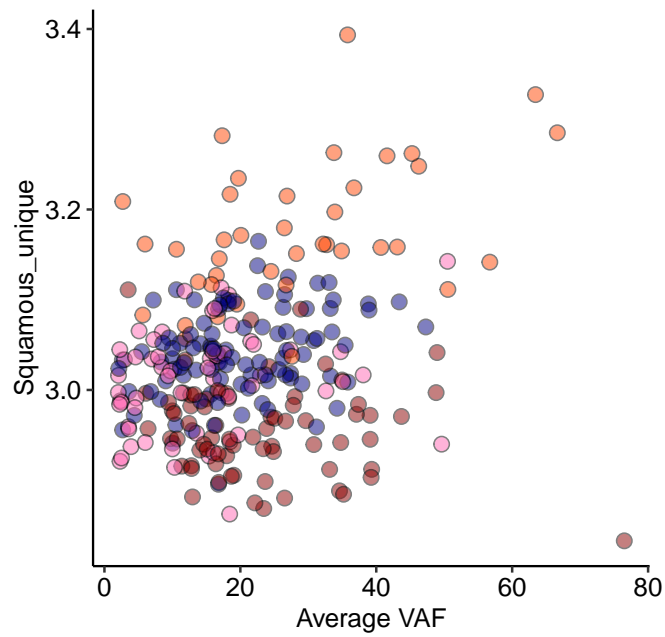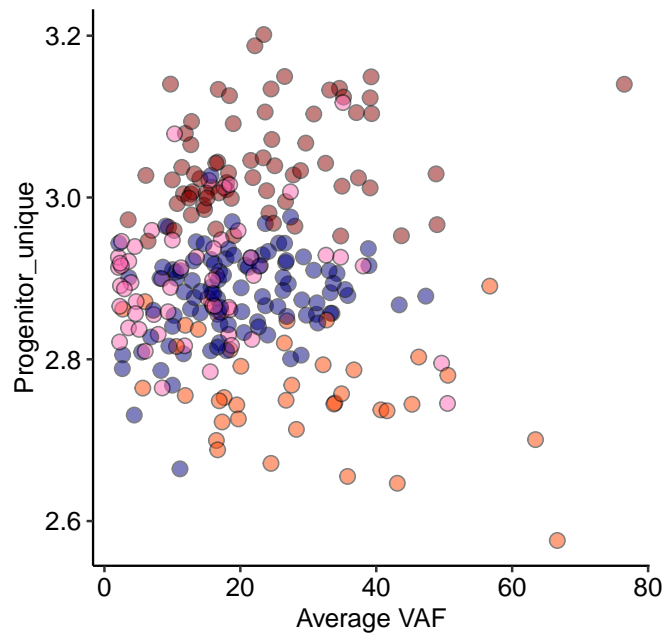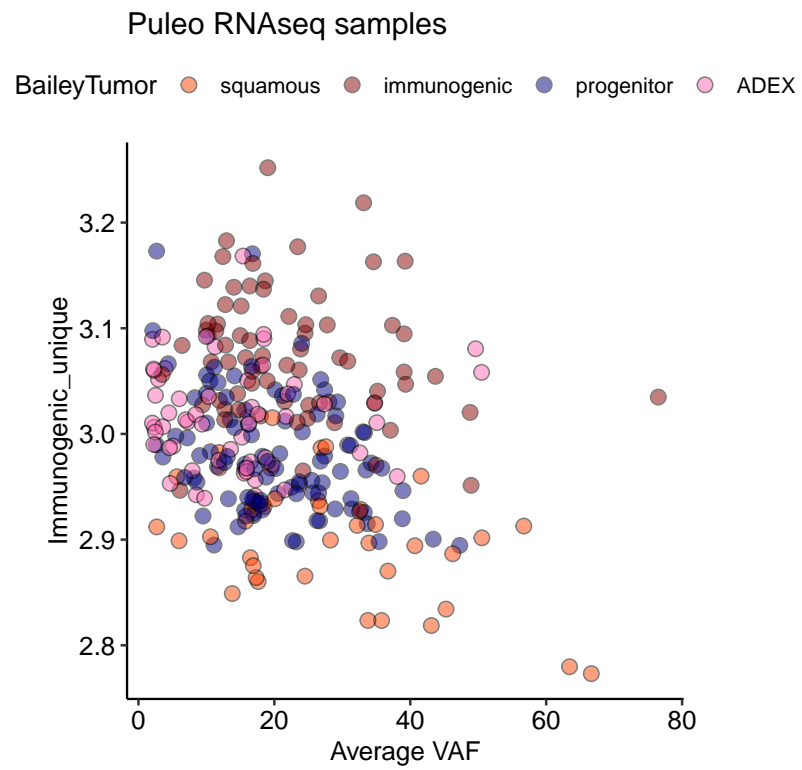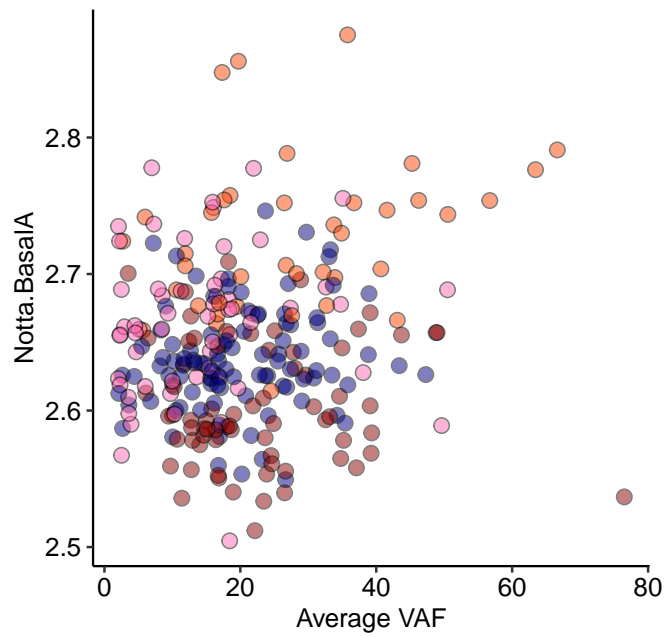
```
## [1] -0.00409
## [1] 0.9468884
```


Puleo RNAseq samples

```
## [1] -0.26057
## [1] 1.555e-05
```

Puleo RNAseq samples
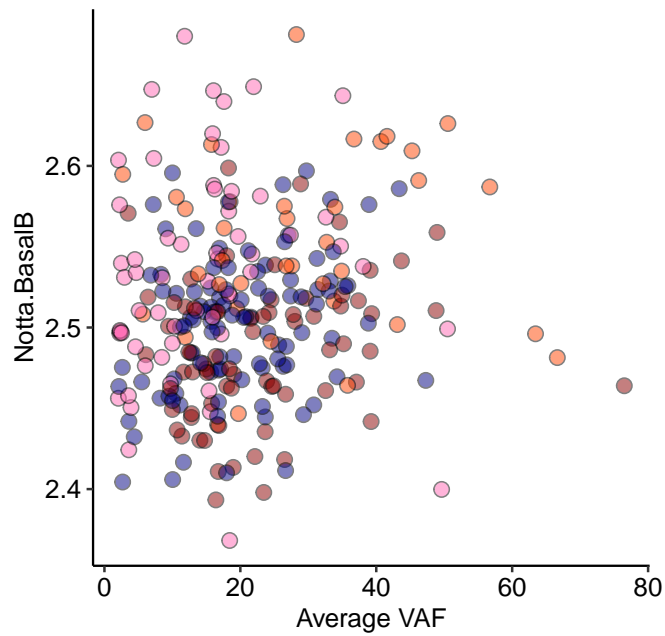
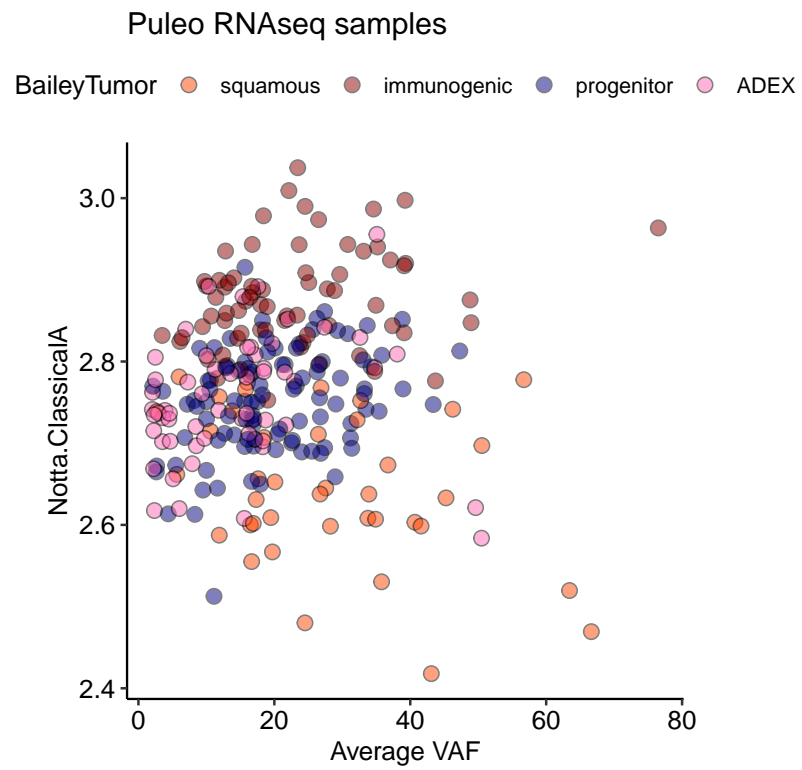BaileyTumor ● squamous ● immunogenic ● progenitor ● ADEX

[1] 0.13884
[1] 0.02300767

Puleo RNAseq samples

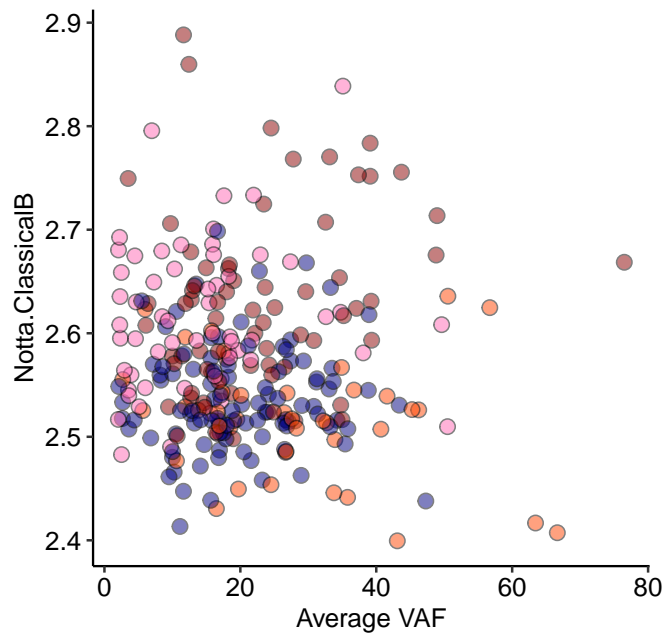BaileyTumor ● squamous ● immunogenic ● progenitor ● ADEX

```
## [1] 0.1301
## [1] 0.03326556
```

Puleo RNAseq samples


```
## [1] 0.02893
## [1] 0.6373474
```

Puleo RNAseq samples


```
## [1] -0.02574
## [1] 0.6748186
```
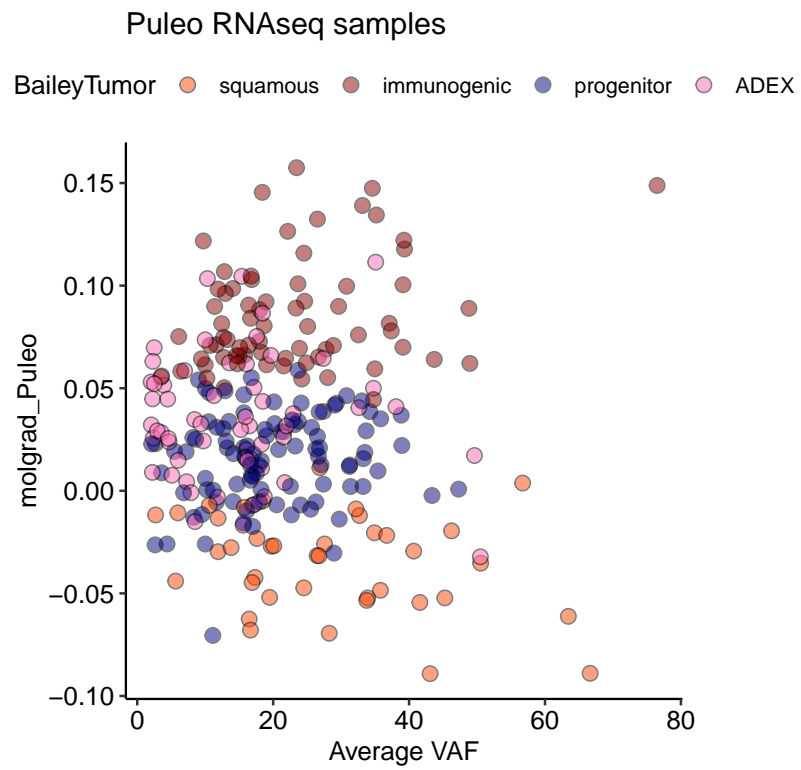
Puleo RNAseq samples

```
## [1] 0.00755
## [1] 0.9021067
```

Puleo RNAseq samples

BaileyTumor ● squamous ● immunogenic ● progenitor ● ADEX


```
## [1] -0.0429
## [1] 0.4843792
```

Puleo RNAseq samples


```
## [1] -0.07827
## [1] 0.2014654
```

Puleo RNAseq samples

```
## [1] 0.13805
## [1] 0.02380719
```

Puleo RNAseq samples

BaileyTumor ● squamous ● immunogenic ● progenitor ● ADEX


```
## [1] 0.25157
## [1] 3.095e-05
```

Puleo RNAseq samples

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## [1] 0.00755
## [1] 0.9021067
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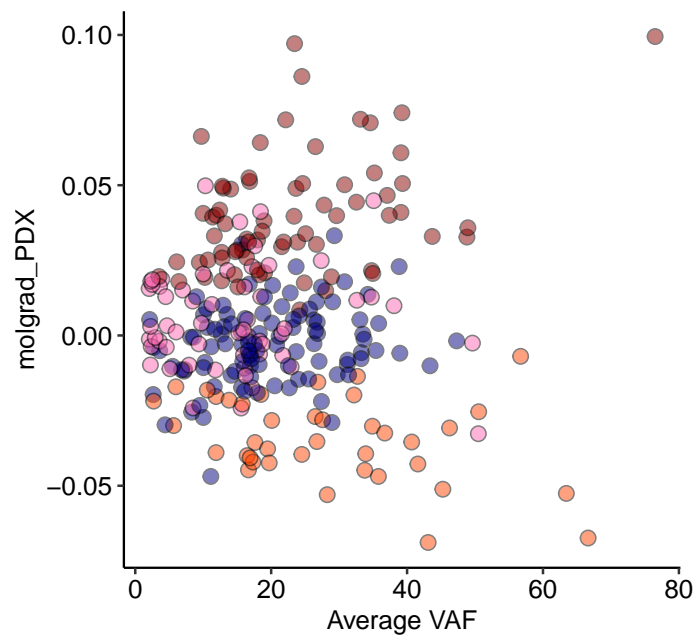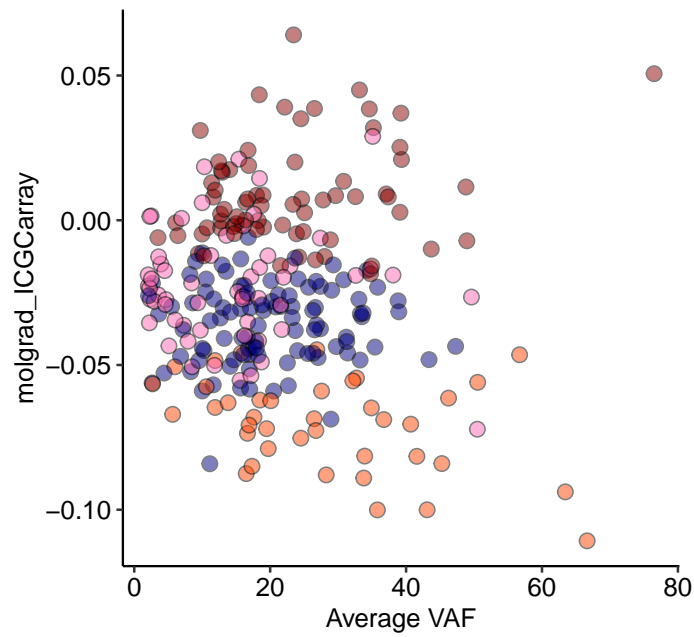
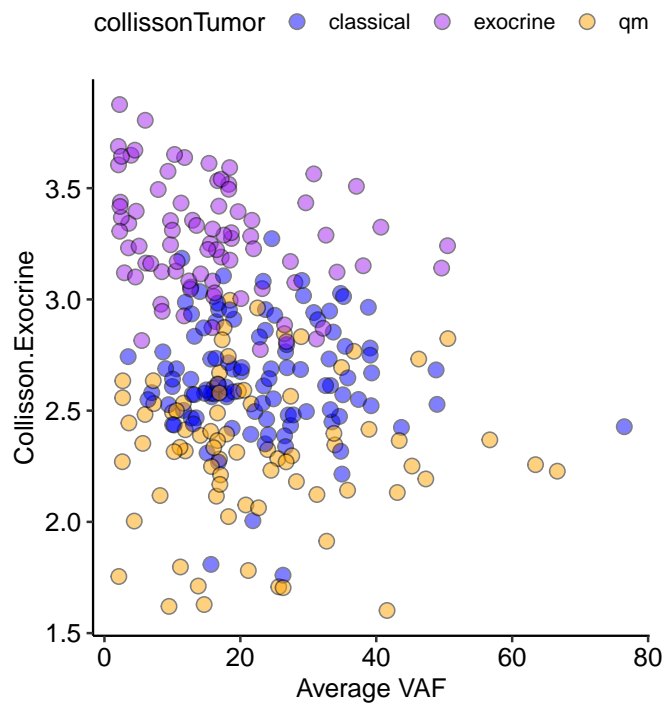**Signatures and average VAF colored by Collisson subtypes**

Puleo RNAseq samples

[1] -0.27207
[1] 6.22e-06

Puleo RNAseq samples


```
## [1] -0.27956
## [1] 3.35e-06
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Puleo RNAseq samples

```
## [1] -0.23084
## [1] 0.00013729
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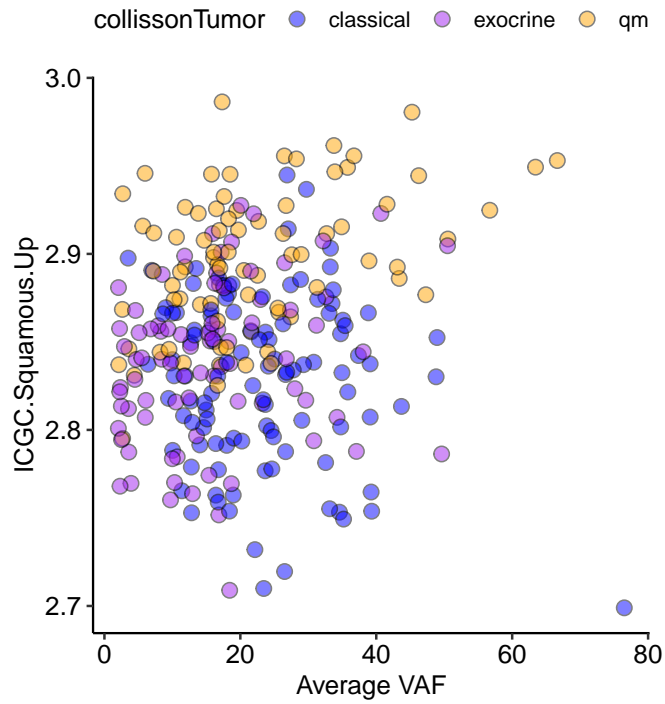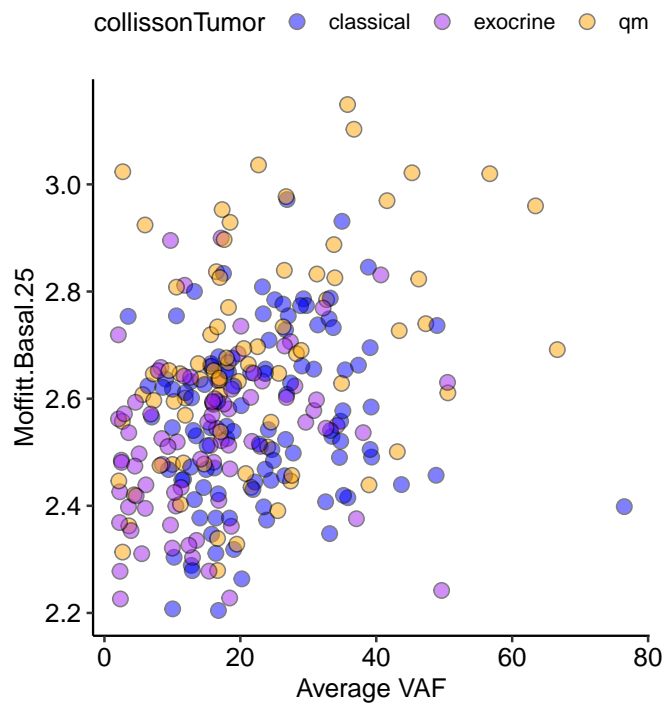
Puleo RNAseq samples


```
## [1] 0.32705
## [1] 4e-08
```

Puleo RNAseq samples

collissonTumor ● classical ● exocrine ● qm


```
## [1] 0.13103
## [1] 0.03201589
```

Puleo RNAseq samples

collissonTumor ● classical ● exocrine ● qm

```
## [1] 0.28846
## [1] 1.56e-06
```

Puleo RNAseq samples


```
## [1] -0.22488
## [1] 0.00020575
```

Puleo RNAseq samples

collissonTumor ● classical ● exocrine ● qm


```
## [1] 0.24632
## [1] 4.571e-05
```

Puleo RNAseq samples

collissonTumor ● classical ● exocrine ● qm

```
## [1] -0.00409
## [1] 0.9468884
```

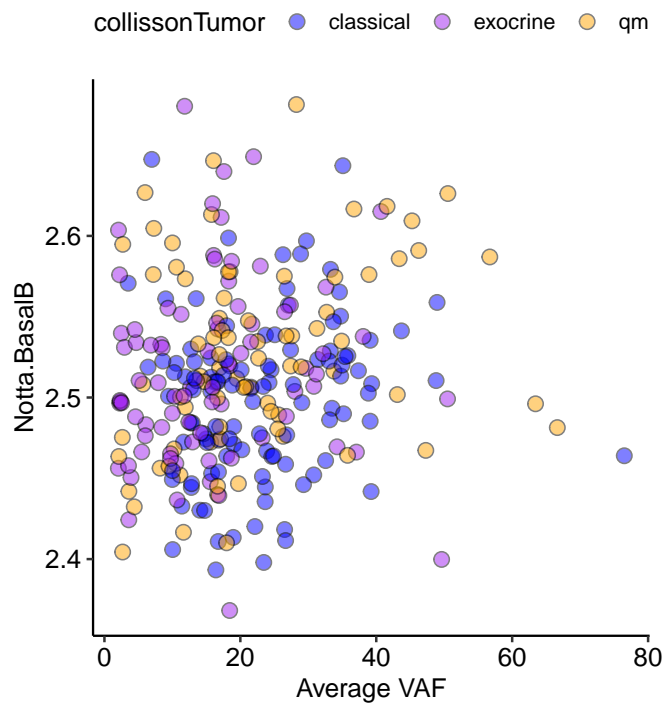## Puleo RNAseq samples

collissonTumor   ● classical   ● exocrine   ● qm
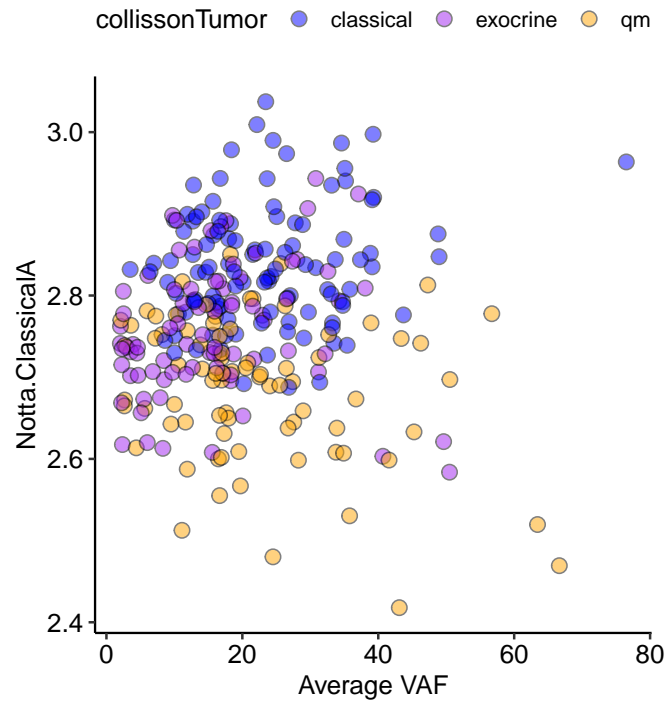


```
## [1] -0.26057
## [1] 1.555e-05
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Puleo RNAseq samples

[1] 0.13884
[1] 0.02300767

Puleo RNAseq samples

```
## [1] 0.1301
## [1] 0.03326556
```
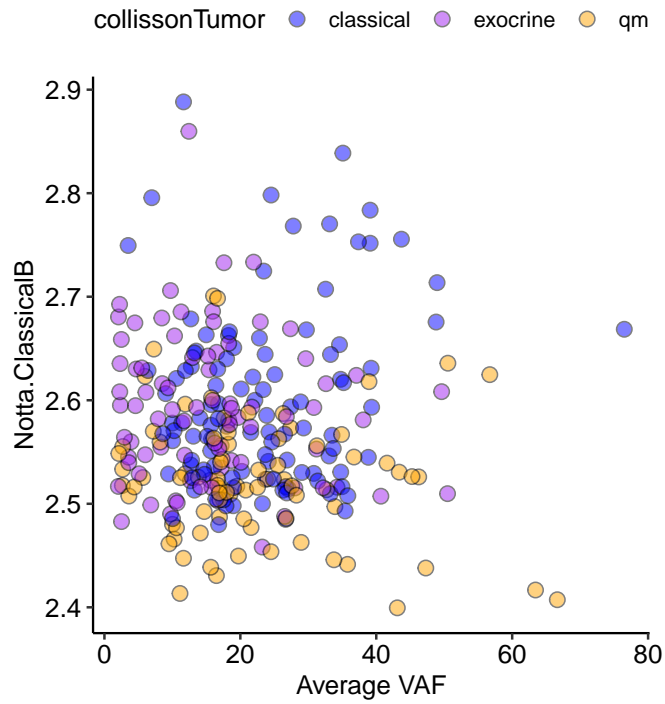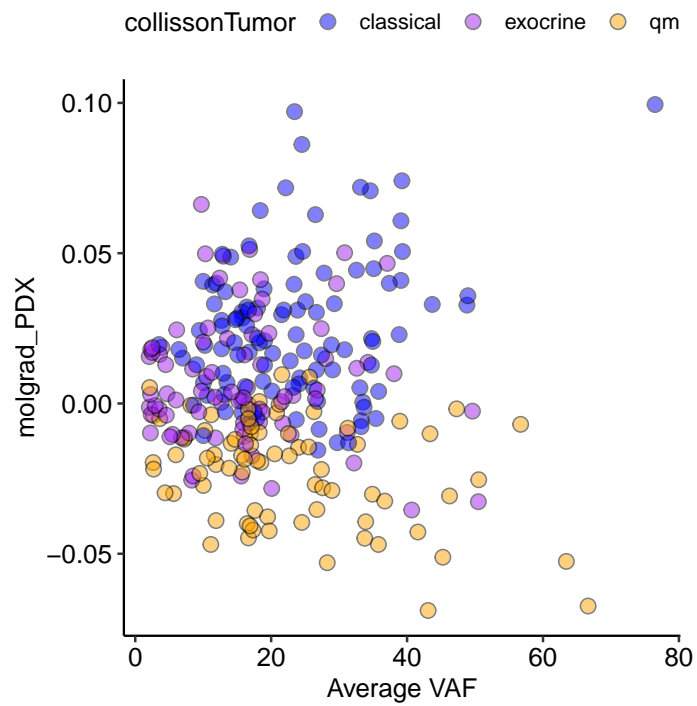
Puleo RNAseq samples


```
## [1] 0.02893
## [1] 0.6373474
```
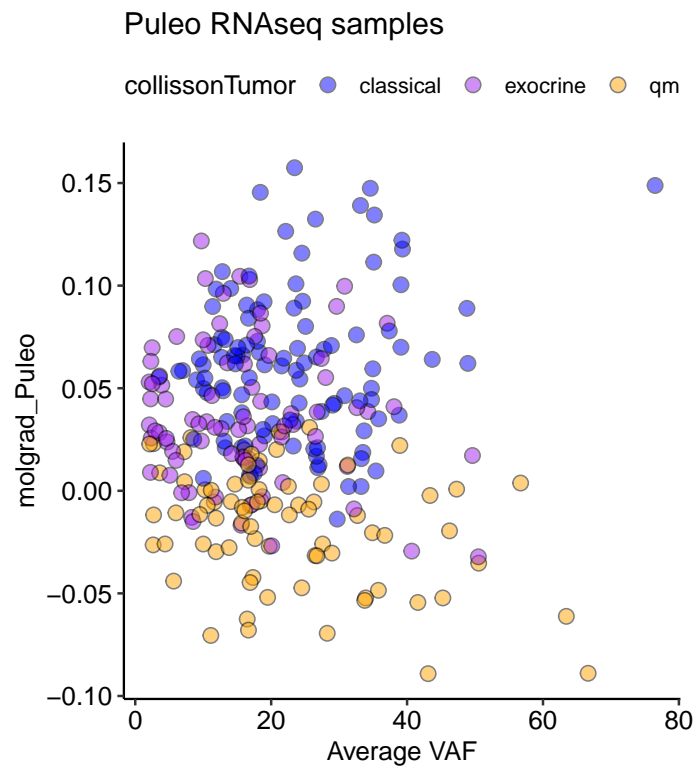
Puleo RNAseq samples
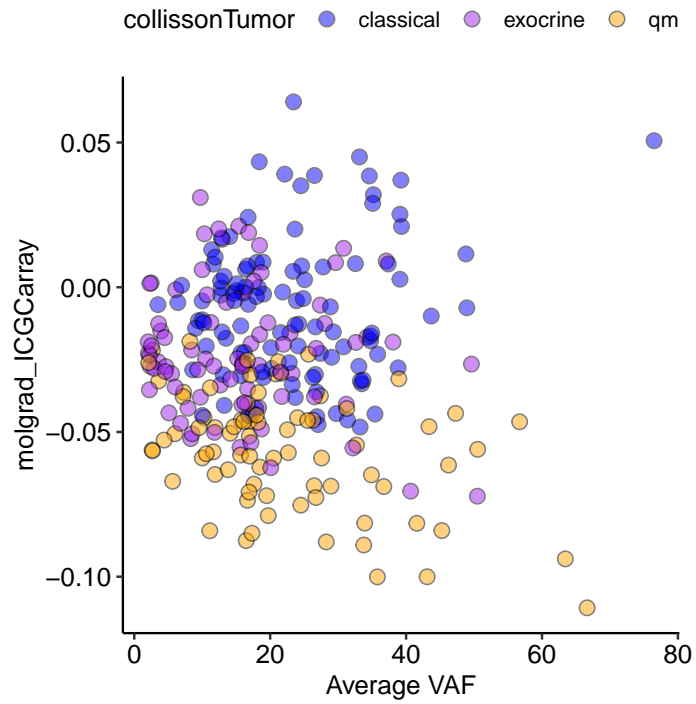
[1] -0.02574
[1] 0.6748186

Puleo RNAseq samples

```
## [1] 0.00755
## [1] 0.9021067
```
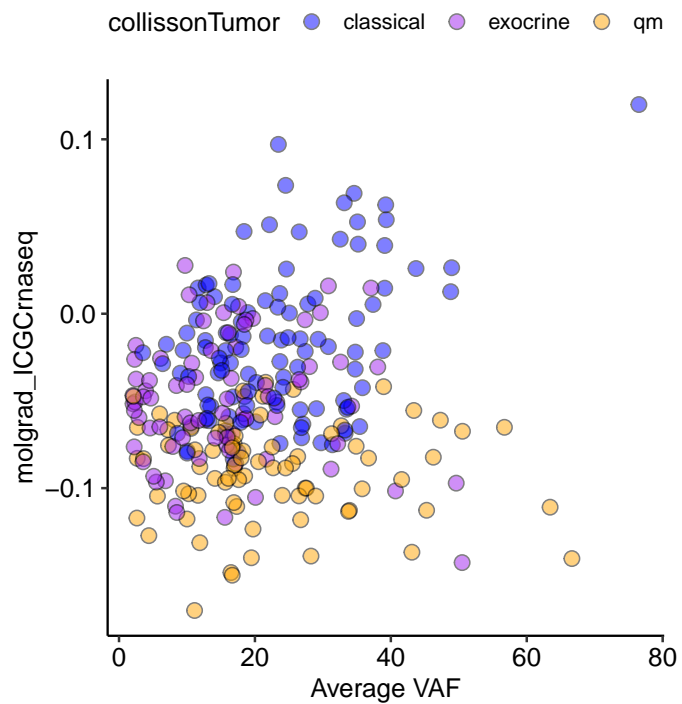
Puleo RNAseq samples


```
## [1] -0.0429
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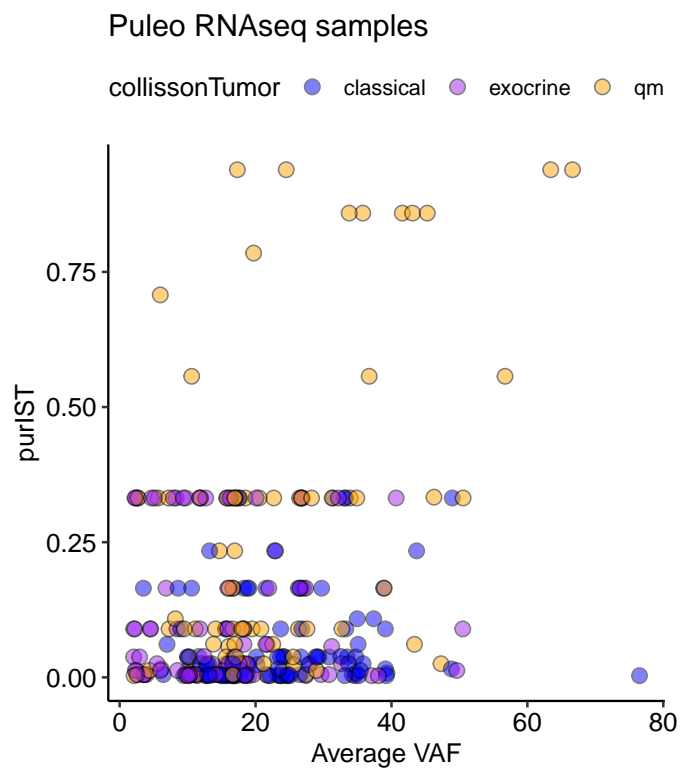
Puleo RNAseq samples

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Puleo RNAseq samples

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## [1] 0.13805
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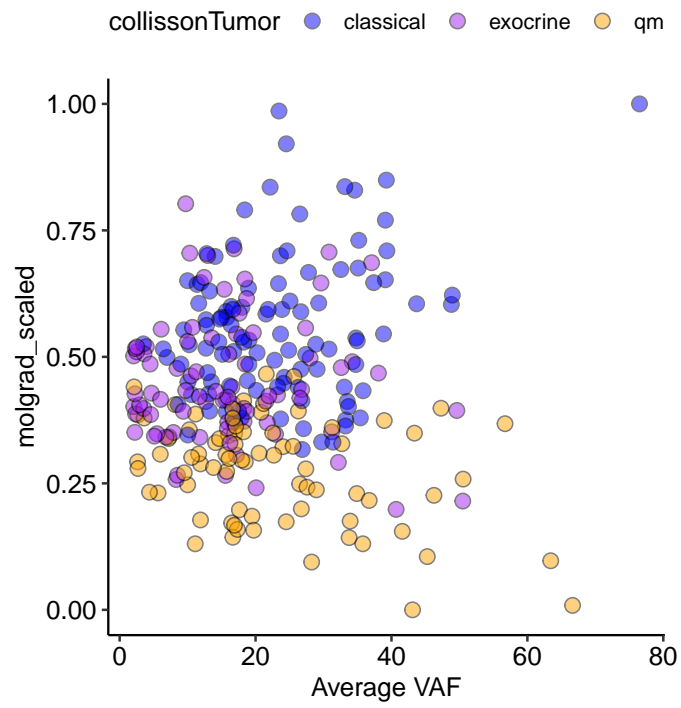
Puleo RNAseq samples


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Puleo RNAseq samples

collissonTumor ● classical ● exocrine ● qm


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## [1] 0.00755
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Signatures and average VAF colored by Moffitt subtypes
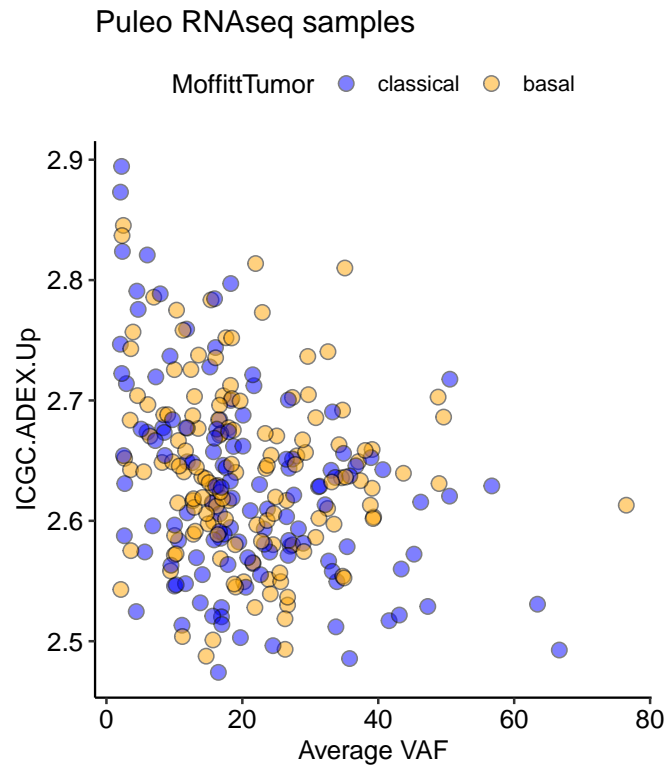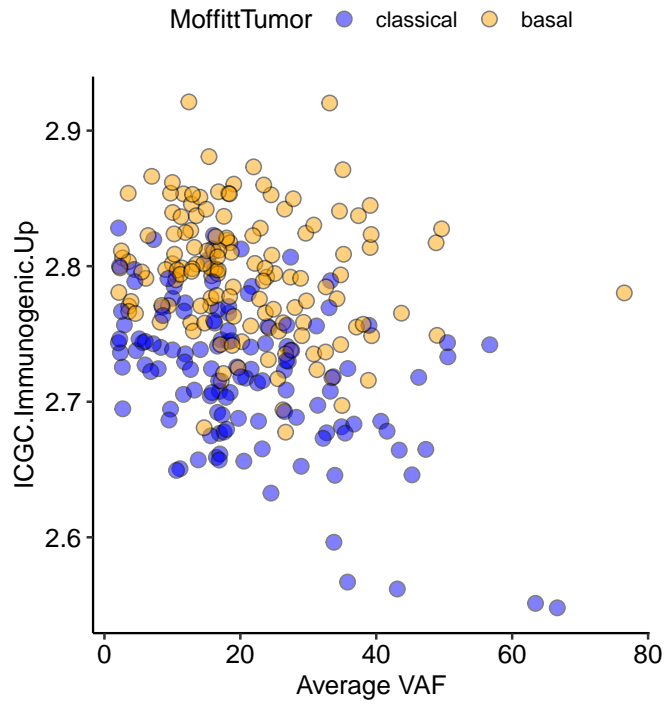
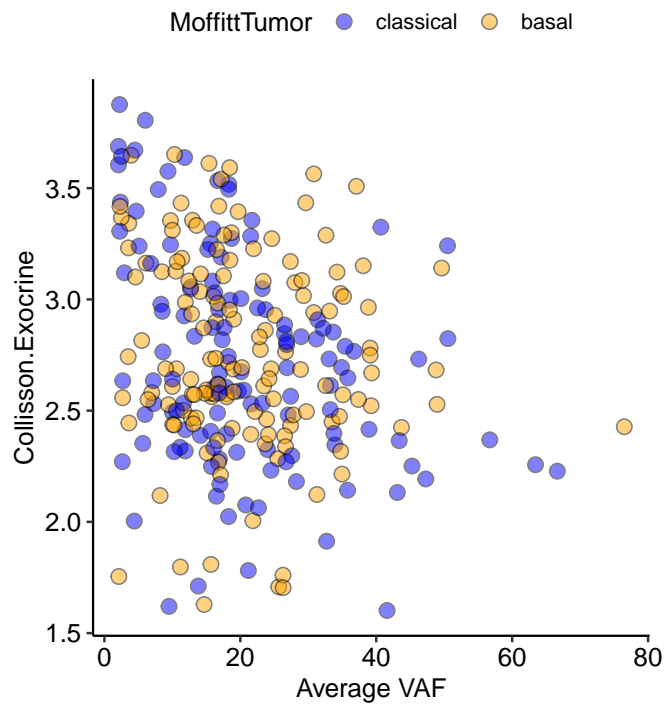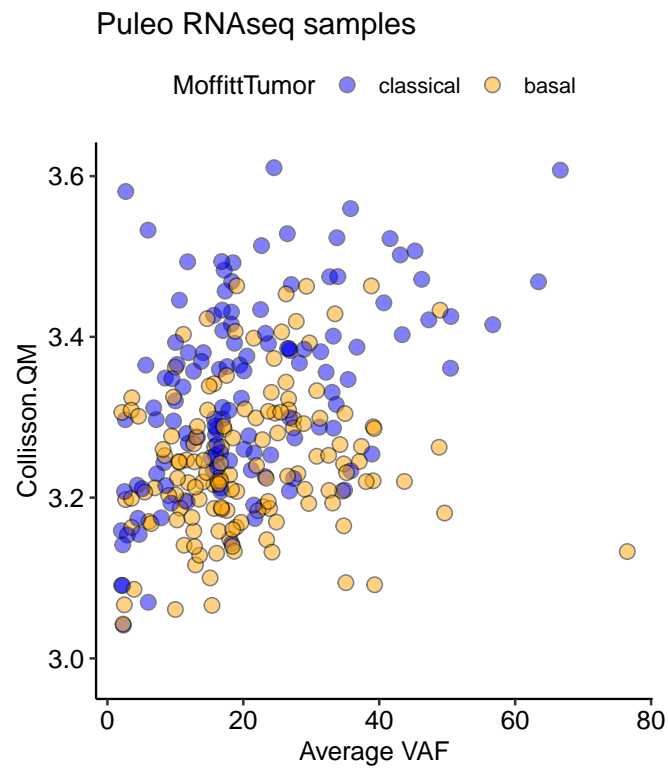## Puleo RNAseq samples


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## [1] -0.27207
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Puleo RNAseq samples


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## [1] -0.27956
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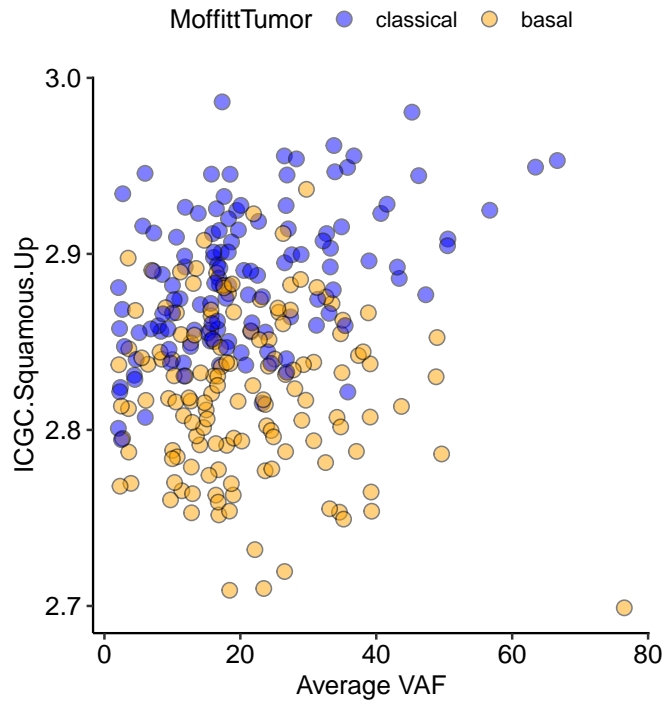
Puleo RNAseq samples

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## [1] -0.23084
## [1] 0.00013729
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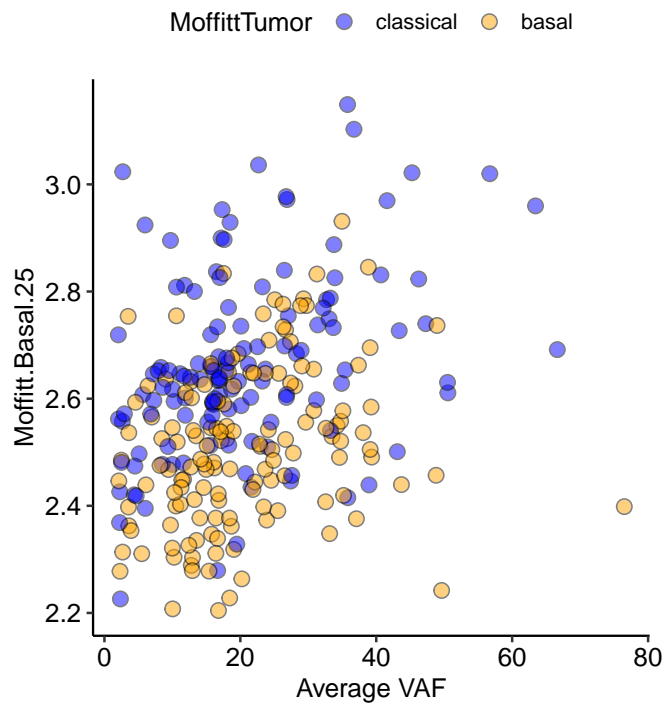
Puleo RNAseq samples


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## [1] 0.32705
## [1] 4e-08
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Puleo RNAseq samples


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## [1] 0.13103
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Puleo RNAseq samples

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## [1] 0.28846
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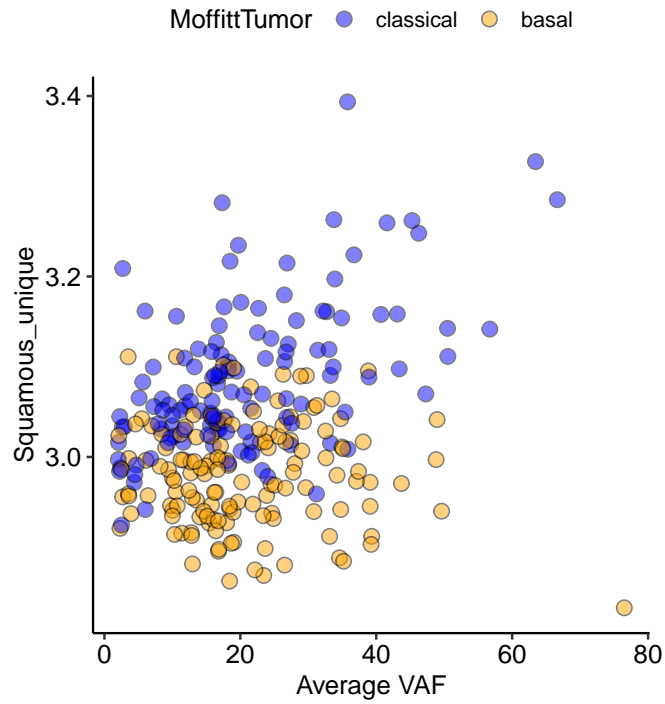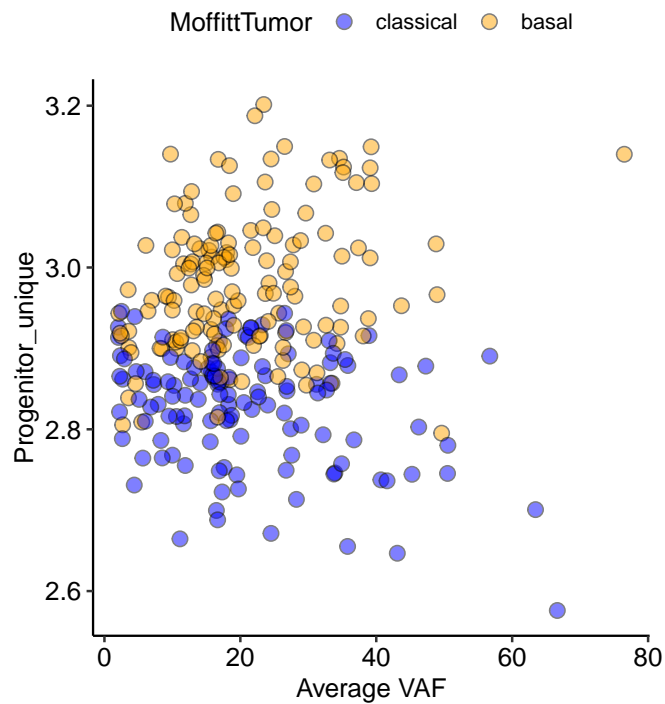
Puleo RNAseq samples


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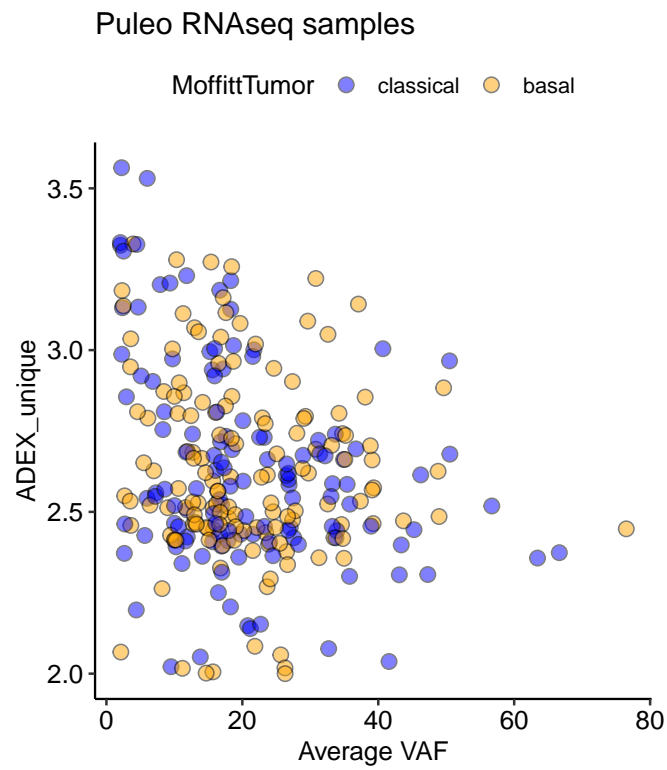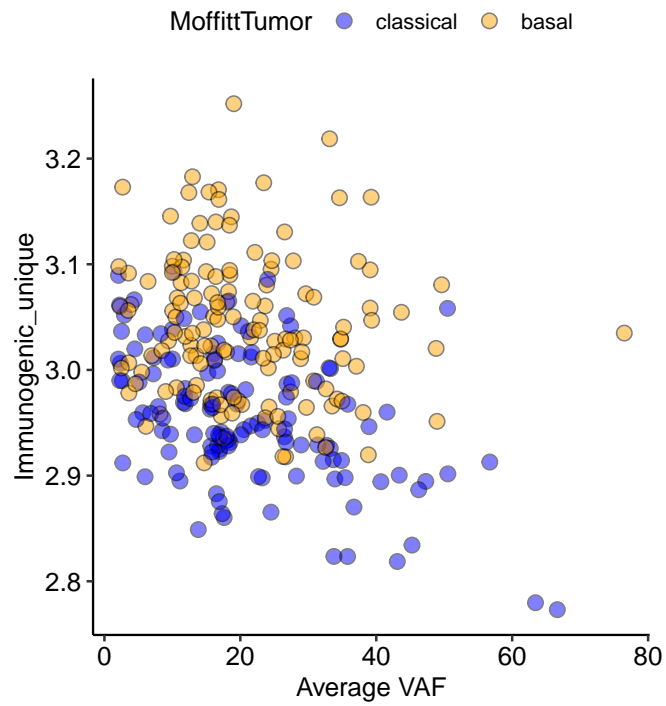
Puleo RNAseq samples


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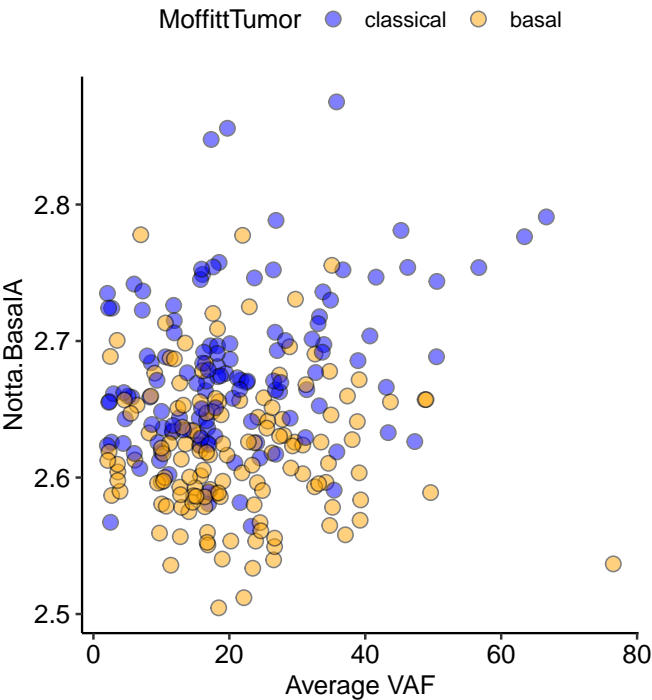
Puleo RNAseq samples

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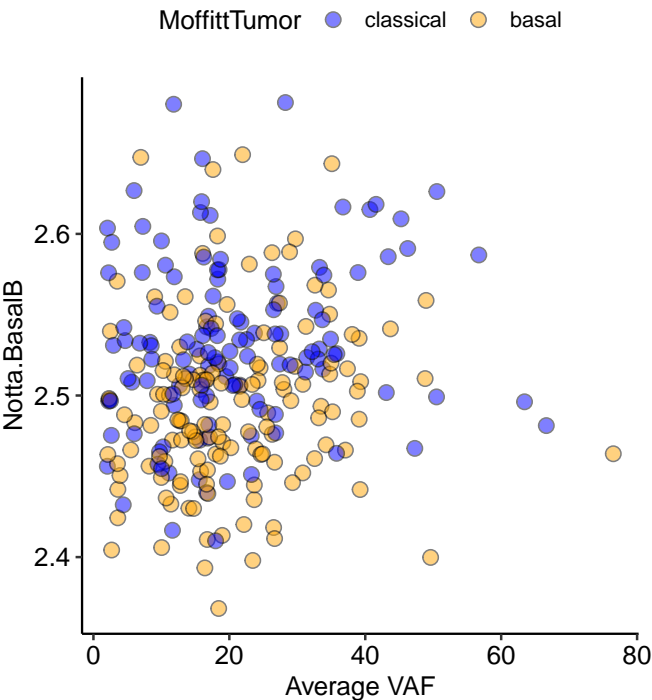
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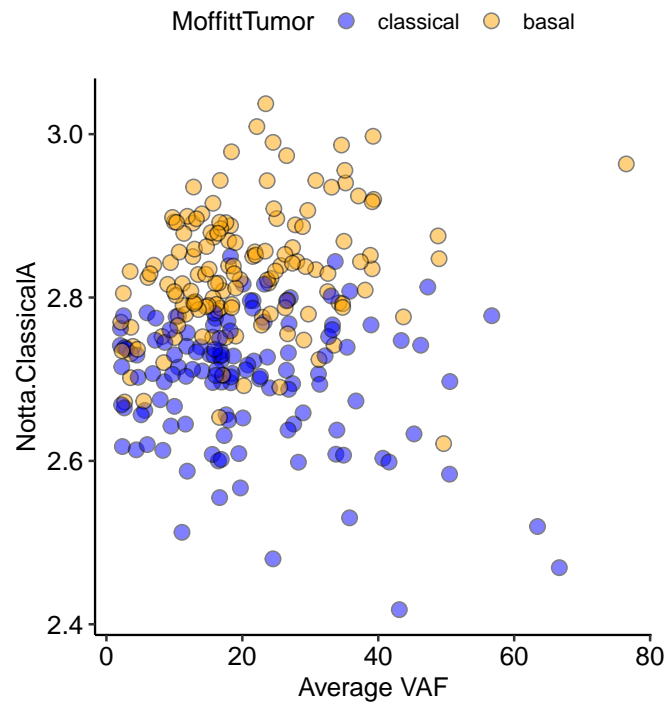Puleo RNAseq samples

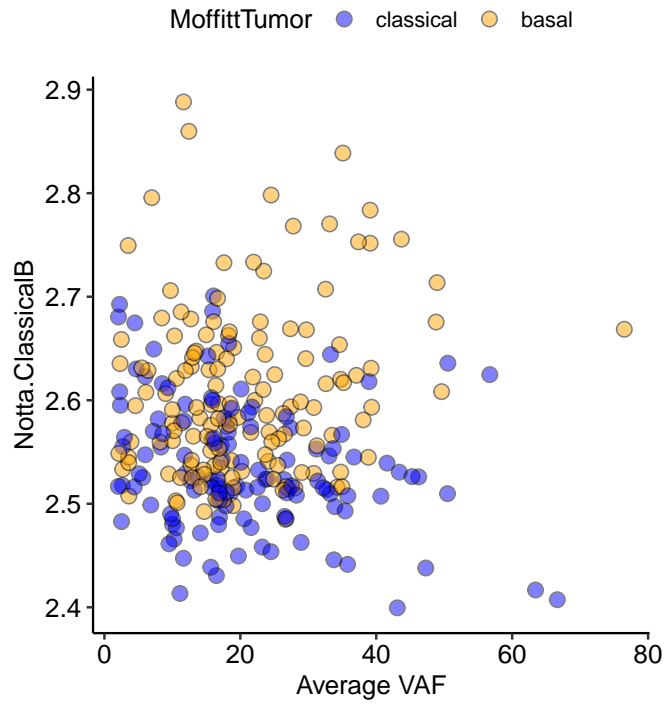## [1] 0.13884
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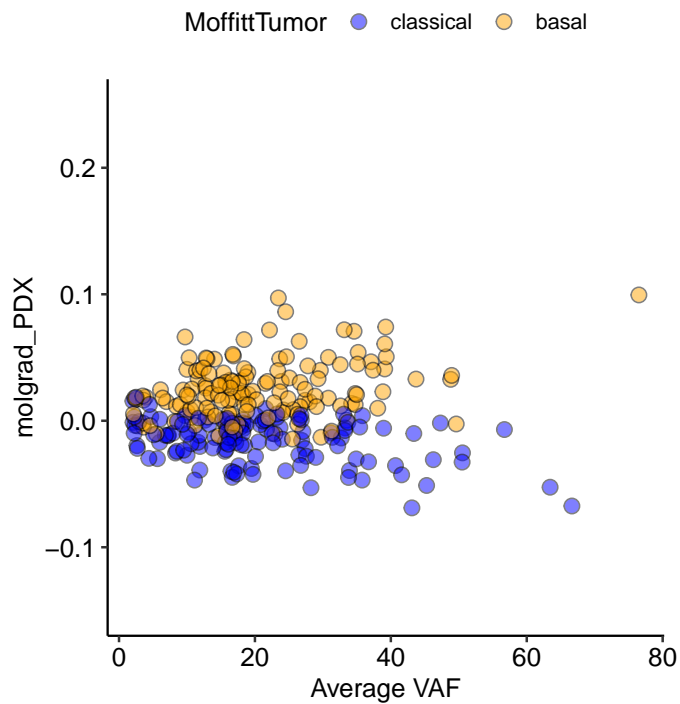
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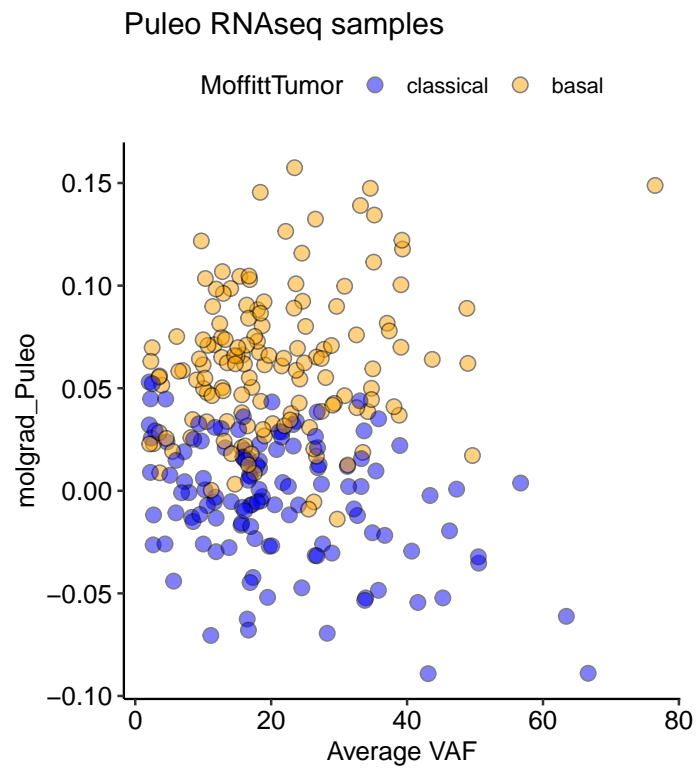
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Puleo RNAseq samples
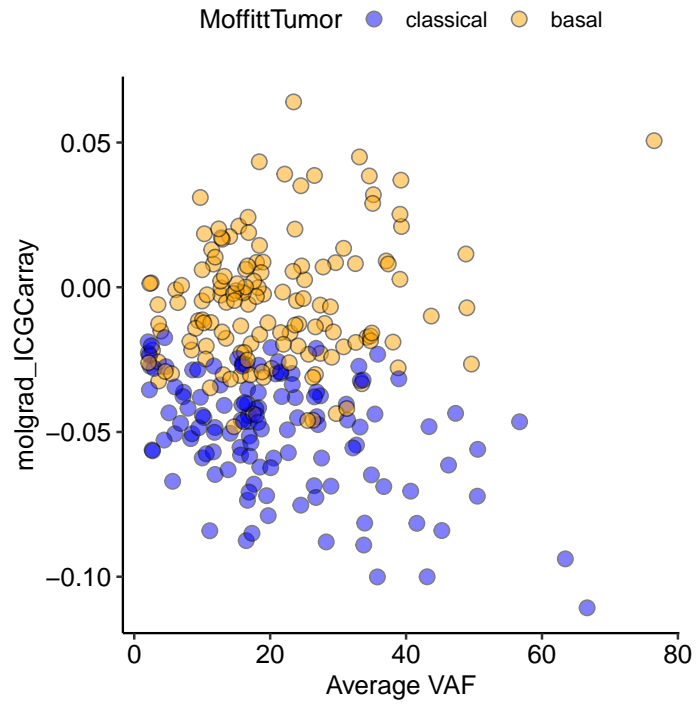
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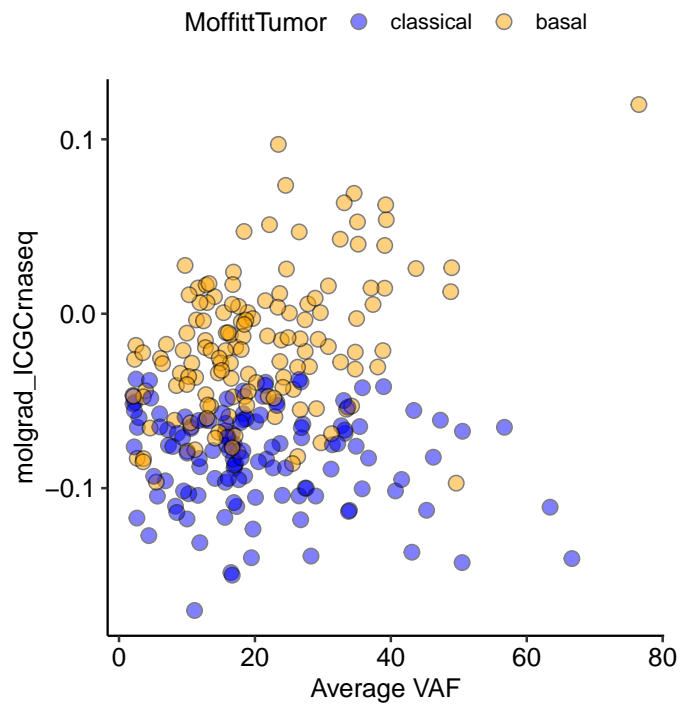
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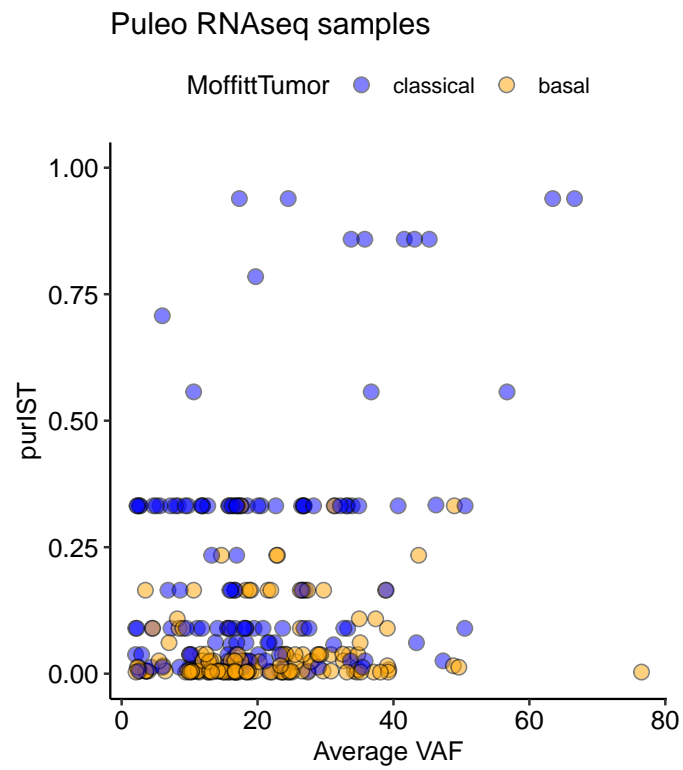
Puleo RNAseq samples

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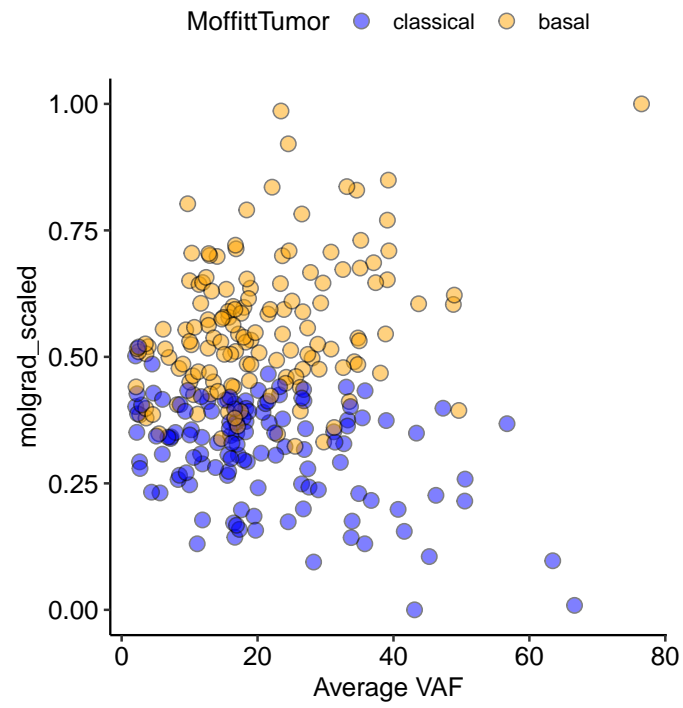## Puleo RNAseq samples


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