

# **Have Env – Will Travel**

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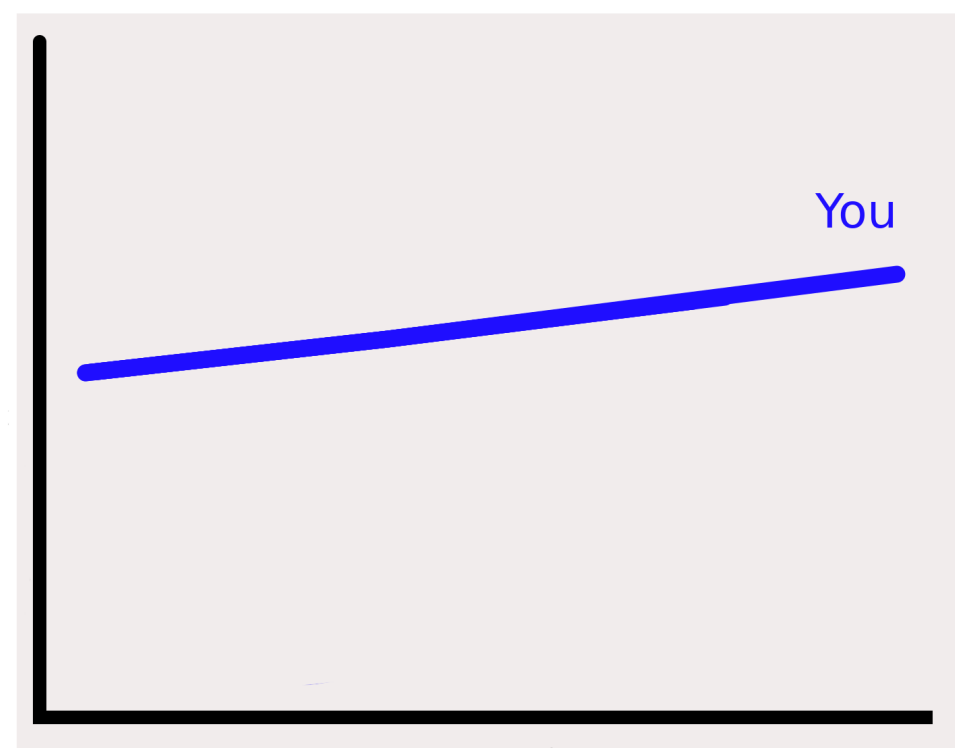
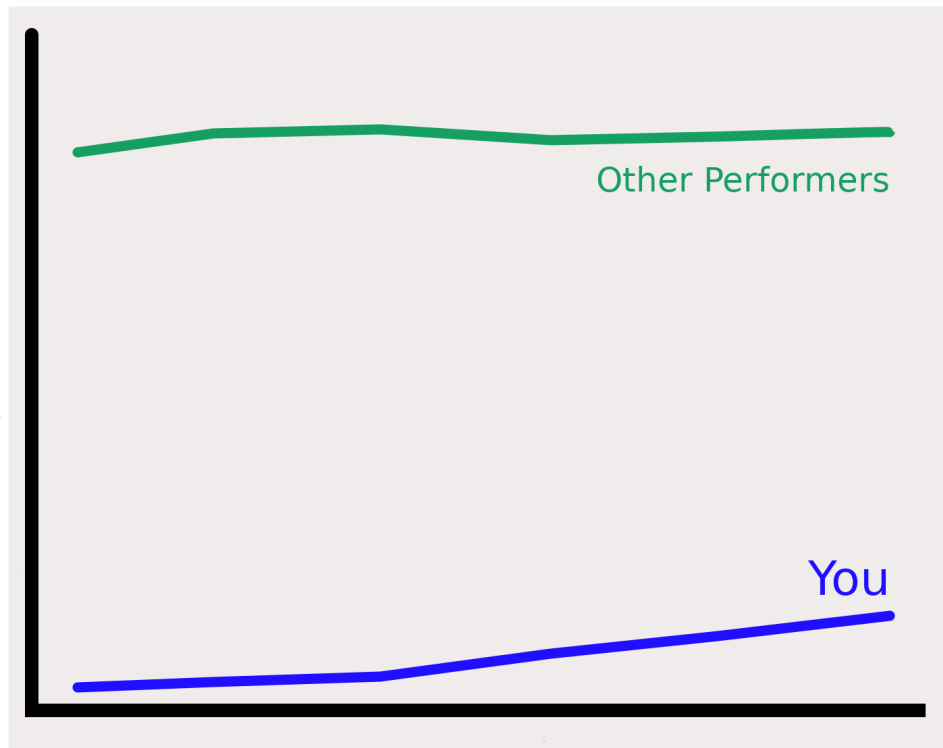
# Tailoring Performance Feedback

**One report does not fit all performers.**

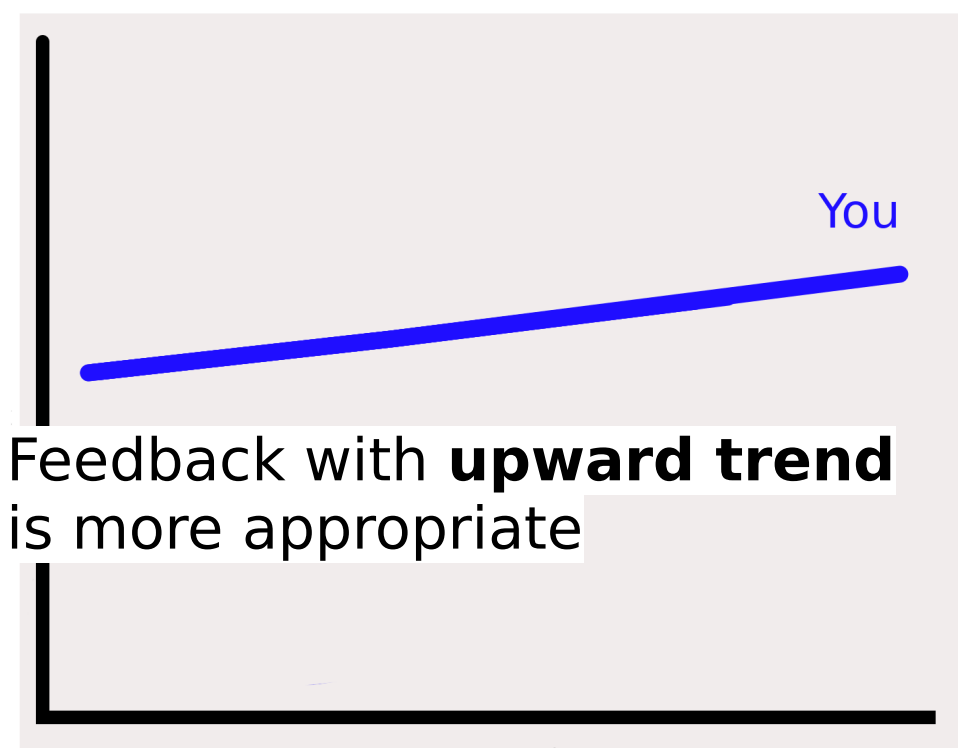
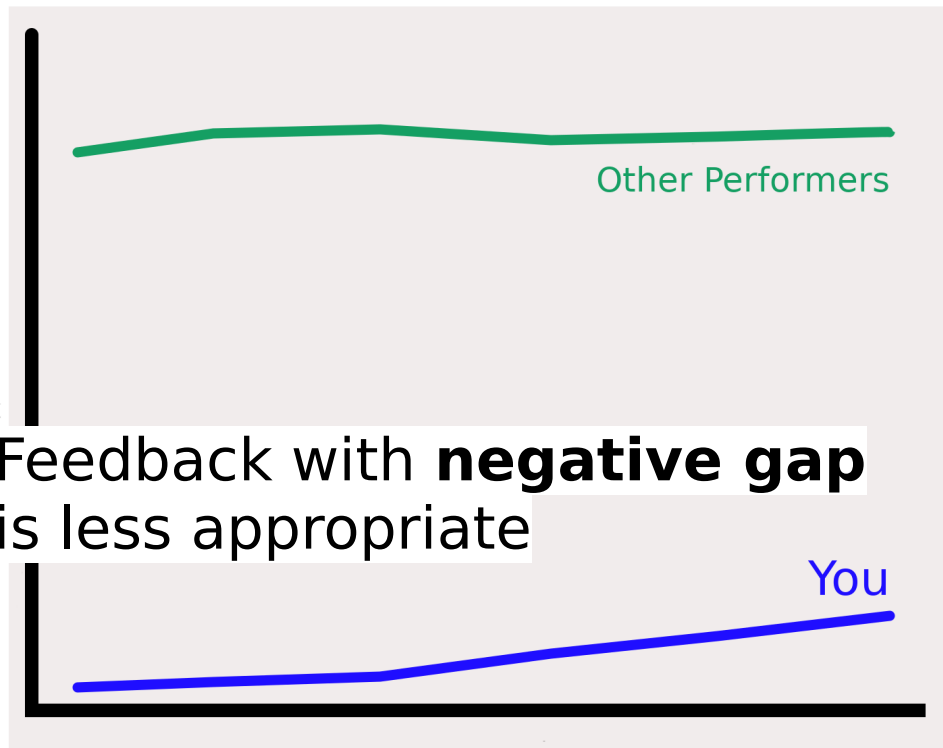
**Psychology has theories about what works and when.**

**Can analyze performance data to select better feedback.**

# E.g Feedback for a beginner



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# Performer Attributes

**Tailoring depends on attributes like:**

- negative\_gap**
- upward\_trend**

**Specific to and specified by client**

*“An upward trend is when someone’s last three counts are increasing.”*

*“A negative gap is when the last count is less than 90% of the average.”*

# Example Data

id	count
Alice	4
Bob	6
Carol	8

# Getting Attributes from Data

*“A negative gap is when the last count is less than 90% of the average.”*

```
eval_gap(count, mean){  
  ifelse( count < mean * 0.9, TRUE, FALSE ) }
```

```
annotate_perf_gap_neg <- function(data){  
  data %>%  
    mutate(mean = mean( count ),  
           perf_gap_neg = eval_gap(count, mean)) %>%  
    select(id, perf_gap_neg) }
```

# Example Annotation

id	perf_gap_neg
Alice	TRUE
Bob	FALSE
Carol	FALSE



# Handling Annotations

**Where to keep the functions.**

**A naming convention for annotations.**

**Standard parameters.**

**Reading in annotation functions.**

**A reasonable way to execute the functions.**

# Where to Keep Annotations

Keep along side the data and config for a client project.

```
/path/to/clients/  
└─ baz-clinic  
    └─ 2018-anti-microbial-stewardship  
        └─ annotations.r  
        └─ config.json  
        └─ performance-data.csv
```

# Annotation Interface

## **Function naming convention:**

`annotate_attribute_label`

e.g. `annotate_upward_trend`

## **Standard parameters:**

`function(performance_data, column_specification)`

# Sourcing Annotations

Source annotations into a new environment:

```
source_annotations <- function(path) {  
  anno_env <- new.env(parent = .BaseNamespaceEnv)  
  source(path, local = anno_env)  
  return(anno_env)  
}
```

local     TRUE, FALSE or **an environment, determining where the parsed expressions are evaluated.**

# Executing Annotations

**Get a list of the functions.**

**Send the data to each of them.**

**Compile results.**

# Get List of Functions

Environment is a object on which we can run:

```
anno_func_names <-
```

```
  lsf.str(envir = anno_env, pattern = "annotate")
```

## List Objects and their Structure

lsf.str return an object of class "ls\_str", basically the **character vector of matching names (functions only for lsf.str)**

# Pass Data to Functions

```
# Build argument list to pass to each of the  
annotation functions.
```

```
aargs <- list(data = data, col_spec = col_spec)
```

```
# Collect table per each annotation function
```

```
anno_results <-  
  lapply(anno_func_names,  
         do.call,  
         args = anno_args,  
         envir = anno_env)
```

# Compile Results

# Reduce results list into a single table

**Reduce**(left\_join, anno\_results)

id	perf_gap_neg	upward_trend	capability_barrier
Alice	TRUE	TRUE	TRUE
Bob	FALSE	TRUE	FALSE
Carol	FALSE	FALSE	FALSE



# Nice Things About Environments

- Don't pollute RGlobal\_Env namespace.
- Can be passed around as a single object.
- Can be mocked for testing.

# Building Envs for Testing

```
bad_env <- new.env()  
bad_env$foo <- function(){}  

```

```
test_that('Annotate complains about no annotation functions.', {  
  expect_success(expect_warning(  
    annotate(mock_data, bad_env, col_spec)  
  ))  
})
```

# Non-Polluting\* Environment

```
env_foo<- new.env()  
env_foo$bar <- function(){print("baz")}
```

```
bar()  
#> Error in bar() : could not find function "bar"
```

```
env_foo$bar()  
#> [1] "baz"
```

\* Calls to library() or require() in annotations.r will modify search path

# Alternatives

- Build packages of annotations
- Source directly into Rglobal\_Env
- Have a separate branch in git for every client
- “How about a web page app that writes the settings that describes the math for the annotations?”

End of Line