

QTIFS

2025-11-05

This analysis uses data compiled from a comprehensive literature review of publications addressing queer and trans fieldwork safety in ecology and related disciplines. The dataset (`QTIFS lit review data Nov 2025.xlsx`) includes 61 unique articles identified through structured searches and screening.

Each article was coded for one or more recommendations to improve field safety, categorized by theme (e.g., Climate, Protocols, Training, Accessibility) and by implementation scale (e.g., Individual, Lab, PI, Department, Institution, Outside Institution). Additional coding captured whether authors described a specific plan for implementation or provided evidence of real-world testing.

The following analyses summarize the distribution of these recommendations, visualize their thematic and structural patterns, and quantify the degree to which certain categories and scales are over- or underrepresented among actionable implementation plans.

INTRO

Attach packages

```
library(tidyverse)    # includes dplyr, ggplot2, readr, forcats, etc.
library(readxl)        # for reading Excel files if used elsewhere
library(janitor)       # for data cleaning and column name formatting
library(ggpubr)         # for ggplot helpers and statistical annotations
library(patchwork)     # for combining multiple ggplots
library(viridis)        # for color palettes
library(ggalluvial)     # for alluvial/flow diagrams
library(scales)         # for axis scaling and labels
```

Load Data

```
q <- read_xlsx("QTIFS lit review data Nov 2025.xlsx") %>%
  janitor::clean_names() %>%
  dplyr::filter(!is.na(recommendation_categories) & !is.na(recommendation_scale)) %>%
  mutate(
    recommendation_categories = fct_relevel(recommendation_categories,
                                              c("Climate",
                                                "Protocols",
                                                "Training",
                                                "Accessibility")),
    recommendation_scale = fct_relevel(recommendation_scale, c(
      "Outside Institution",
      "Institution",
      "Department",
```

```

    "Lab", "PI",
    "Individual"
  ))
)

arts <- q %>% dplyr::group_by(as.factor(recommendation_scale)) %>%
  dplyr::summarize(n())

sum(arts$n())
## [1] 165

print(arts)

## # A tibble: 6 x 2
##   `as.factor(recommendation_scale)` `n()`
##   <fct>                <int>
## 1 Outside Institution      18
## 2 Institution             42
## 3 Department              16
## 4 Lab                      11
## 5 PI                       50
## 6 Individual               28

q %>% dplyr::group_by(as.factor(recommendation_scale)) %>% dplyr::summarize(n()/nrow(q))

## # A tibble: 6 x 2
##   `as.factor(recommendation_scale)` `n()/nrow(q)`
##   <fct>                  <dbl>
## 1 Outside Institution     0.109
## 2 Institution            0.255
## 3 Department             0.0970
## 4 Lab                     0.0667
## 5 PI                      0.303
## 6 Individual              0.170

```

Set color palette

```

# define palette
my_colors <- c("#003f5c", "#58508d", "#bc5090", "#ff6361", "#ffa600")

```

Heatmap

Visualize the frequency of recommendations across thematic categories and implementation scales.
 ##### wrangle

```

# summarize number of recommendations per category and scale
q_collect <- q %>%
  select(recommendation_categories, recommendation_scale) %>%
  group_by(recommendation_categories, recommendation_scale, .drop = FALSE) %>%
  dplyr::summarize(
    num = n()
  ) %>%
  ungroup() %>%
  na.omit()

```

`summarise()` has grouped output by 'recommendation_categories'. You can
override using the ` `.groups` argument.

counts

```

# count unique recommendations by category
q_unique_rec <- q %>%
  select(article, one_sentence_summary, recommendation_categories, recommendation_scale) %>%
  distinct() %>%
  count(recommendation_categories, name = "num")

# count unique combinations of category and scale
q_heatmap <- q %>%
  select(article, one_sentence_summary, recommendation_categories, recommendation_scale, recommendation)
distinct() %>%
  count(recommendation_categories, recommendation_scale, name = "num")

# count unique articles and recommendations to use in plots later
q_unique_articles <- length(unique(q$article))

q_unique_recs_num <- q %>%
  select(article, recommendation_categories, recommendation_scale) %>%
  distinct() %>%
  summarize(n())

```

PLOT

```

sum_q <- sum(q_heatmap$num)

heatmap <- ggplot(q_heatmap, aes(x = recommendation_scale, y = recommendation_categories, size = num, f
  annotate("rect", xmin = 5.5, xmax = Inf, ymin = 0, ymax = Inf, fill = 'gray', alpha = 0.4) +
  geom_vline(xintercept = 5.5, color = "gray", linewidth = 0.75, linetype = "dashed") +
  geom_point(pch = 21) +
  theme_bw() +
  theme(axis.text.y = element_text(face = "italic")) +
  scale_fill_viridis(option = "viridis", name = "Number of\nStudies",
                     breaks = c(1, 3, 9, 15, 21)
  ) +
  scale_size_continuous(range = c(1,15),

```

```

        name = "Number of\nStudies",
        breaks = c(1, 3, 9, 15, 21)
    ) +
scale_x_discrete(limits = rev) +
scale_y_discrete(limits = rev) +
theme(axis.line.x = element_line(linewidth = 1.5,
                                 color = "grey50",
                                 arrow = grid::arrow(length = unit(0.4, "cm"),
                                                     ends = "last")),
      axis.text.x = element_text(angle = 45, hjust = 1, face = c("plain", "plain", "plain", "plain", "plain"),
                                 legend.key.size = unit(1, "cm"))
    ) +
guides(fill = guide_legend(), size = guide_legend()) +
labs(title = "Distribution of recommendation categories and scales", x = "Scale", y = "Categories", s
## Warning: Vectorized input to `element_text()` is not officially supported.
## i Results may be unexpected or may change in future versions of ggplot2.

# heatmap
# ggsave("Heatmap.png", width = 7, height = 5)

```

Bar chart

wrangle

```

# define category levels and matching color palette
category_levels <- levels(q_collect$recommendation_categories)

# create a named vector for consistent color mapping
category_palette <- setNames(my_colors[1:length(category_levels)], category_levels)

# summarize total counts for each category
q_category_totals <- q_collect %>%
  group_by(recommendation_categories) %>%
  summarise(total = sum(num)) %>%
  ungroup()

```

PLOT

```

sum_collect <- sum(q_collect$num)

themes <- ggplot(q_collect, aes(x = recommendation_categories, y = num, fill = recommendation_categories))
  geom_bar(stat = "identity") +
  geom_text(
    data = q_category_totals,
    aes(x = recommendation_categories, y = total, label = paste0("N = ", total)),

```

```

    vjust = -1,
    size = 3,
    inherit.aes = FALSE
) +
scale_fill_manual(values = category_palette) +
labs(x = "Recommendation Category", y = "Number of Unique Recommendations",
     title = "Main Themes Across Articles",
     subtitle = paste0("Each article has 1 recommendations; \n N = ", sum_collect , " recommendations ac
ylim(c(0, 90)) +
theme_minimal() +
theme(legend.position = "none",
      panel.grid.major.x = element_blank(),
      axis.text.x = element_text(angle = 45, hjust = 0.75, size = 10),
      panel.background = element_rect(fill = "white", color = NA),
      plot.background = element_rect(fill = "white", color = NA))

# themes
# ggsave("Recommendation Categories Bar Chart.png", width = 6, height = 4)

```

Recommendation plans

Does each article included an explicit plan for implementing recommendations? ##### wrangle

```

# count unique articles with their recommendation plan Y/N
q_plan_status <- q %>%
  filter(!is.na(recommendation_categories) & !is.na(recommendation_scale)) %>%
  select(article, plan_for_implementation_y_n) %>%
  distinct() %>%
  count(plan_for_implementation_y_n, name = "num")

# total counts per plan status
q_plan_totals <- q_plan_status %>%
  group_by(plan_for_implementation_y_n) %>%
  summarise(total = sum(num)) %>%
  ungroup()

# sanity check: check for inconsistencies in plan coding across same articles
inconsistent_plan_check <- q %>%
  select(article, plan_for_implementation_y_n) %>%
  distinct() %>%
  group_by(article) %>%
  summarise(unique_vals = n_distinct(plan_for_implementation_y_n)) %>%
  filter(unique_vals > 1)

# print any inconsistencies: articles that were coded both yes and no
inconsistent_plan_check

## # A tibble: 0 x 2
## # i 2 variables: article <chr>, unique_vals <int>

```

```
q_unique_plans <- sum(q_plan_status$num)
```

PLOT

```
plan <- ggplot(q_plan_status, aes(x = plan_for_implementation_y_n, y = num, fill = plan_for_implementat
  geom_bar(stat = "identity") +
  geom_text(
    data = q_plan_totals,
    aes(x = plan_for_implementation_y_n, y = total, label = paste0("N = ", total)),
    vjust = -1,
    size = 3,
    inherit.aes = FALSE
  ) +
  scale_fill_manual(values = c(my_colors[1], my_colors[3])) +
  labs(
    x = "Stated Plan for Implementation",
    y = "Number of Articles",
    title = "Did Articles Include a Plan for Implementation?",
    subtitle = paste0("N = ", q_unique_plans, " articles")
  ) +
  expand_limits(y = c(0, 60)) +
  scale_x_discrete(breaks = c("N", "Y"), labels = c("No", "Yes")) +
  theme_minimal() +
  theme(
    legend.position = "none",
    panel.grid.major.x = element_blank(),
    axis.text.x = element_text(hjust = 1, size = 10),
    panel.background = element_rect(fill = "white", color = NA),
    plot.background = element_rect(fill = "white", color = NA)
  )

# plan
# ggsave("Plan Inclusion Bar Chart.png", width = 5, height = 5)
```

Real-world testing

Were recommendations explicitly tested or applied in real-world settings?
wrangle

```
# count unique articles that tested recommendations in the real world
q_real_world_test <- q %>%
  select(article, real_worldImplementation_testing_y_n) %>%
  distinct() %>%
  count(real_worldImplementation_testing_y_n, name = "num")

# compute total number of articles per response
q_real_totals <- q_real_world_test %>%
  group_by(real_worldImplementation_testing_y_n) %>%
  summarise(total = sum(num)) %>%
  ungroup()
```

```

# check for inconsistencies in coding for real-world testing
realworld <- q %>% dplyr::select(article, recommendation_categories, recommendation_scale, real_world_implementations)

# identify any citations where multiple values exist
inconsistent_test <- realworld %>%
  group_by(article) %>%
  summarise(unique_vals = n_distinct(real_world_implementations_y_n)) %>%
  filter(unique_vals > 1)

# show the inconsistent citations (if any)
inconsistent_test

## # A tibble: 0 x 2
## # i 2 variables: article <chr>, unique_vals <int>

q_real_world_test <- q %>%
  select(article, real_world_implementations_y_n) %>%
  distinct() %>%
  count(real_world_implementations_y_n, name = "num")

```

PLOT

```

testing <- ggplot(q_real_world_test, aes(x = real_world_implementations_y_n, y = num, fill = real_world_implementations_y_n))
  geom_bar(stat = "identity") +
  geom_text(
    data = q_real_totals,
    aes(x = real_world_implementations_y_n, y = total, label = paste0("N = ", total)),
    vjust = -1,
    size = 3,
    inherit.aes = FALSE
  ) +
  scale_fill_manual(values = c(my_colors[3], my_colors[1])) +
  labs(x = "Real World Testing of Recommendations", y = "Number of Articles",
       title = "Were Recommendations Tested in the Real-World?",
       subtitle = paste0("N = ", q_unique_articles, " articles")) +
  expand_limits(y = c(0, 50)) +
  scale_x_discrete(breaks = c("N", "Y"), labels = c("No", "Yes")) +
  theme_minimal() +
  theme(legend.position = "none",
        panel.grid.major.x = element_blank(),
        axis.text.x = element_text(hjust = 1, size = 10),
        panel.background = element_rect(fill = "white", color = NA),
        plot.background = element_rect(fill = "white", color = NA))

# testing
# ggsave("Real World Testing Bar Chart.png", width = 5, height = 5)

```

Implementation Responsibility

Identify where the literature places responsibility for implementing recommendations.
wrangle

```
# count how many recommendations were assigned to each implementation scale
q_responsibility <- q %>%
  select(article, recommendation_categories, recommendation_scale) %>%
  distinct() %>%
  count(recommendation_scale, name = "num")

sum(q_responsibility$num)

## [1] 165

q %>% group_by(article, recommendation_scale) %>% dplyr::summarize(n())

## `summarise()` has grouped output by 'article'. You can override using the
## `.groups` argument.

## # A tibble: 146 x 3
## # Groups:   article [61]
##   article           recommendation_scale `n()`
##   <chr>              <fct>                <int>
## 1 Atchinson 2021    Outside Institution     1
## 2 Atchinson 2021    Institution          1
## 3 Baker 2000        Outside Institution     1
## 4 Baker 2000        Institution          1
## 5 Beltran et al 2021 Institution          1
## 6 Beltran et al 2021 Department          1
## 7 Blonder 2022      PI                  3
## 8 Bowser & Cid 2021 Institution          1
## 9 Bowser & Cid 2021 PI                  1
## 10 Bracken and Mawdsley 2004 Individual    1
## # i 136 more rows
```

PLOT

```
pal <- colorRampPalette(my_colors[1:5])

responsibility <- ggplot(q_responsibility, aes(x = recommendation_scale, y = num, fill = recommendation_
  geom_bar(stat = "identity") +
  geom_text(
    aes(x = recommendation_scale, y = num, label = paste0("N = ", num)), vjust = -1, size = 3) +
  scale_fill_manual(values = pal(8)) +
  labs(x = "Scale of Recommendations", y = "Number of Articles",
       title = "Where Should Implementation Fall?", subtitle = paste0("N = ", q_unique_articles, " articles")) +
  expand_limits(y = c(0, 60)) +
  theme_minimal() +
```

```

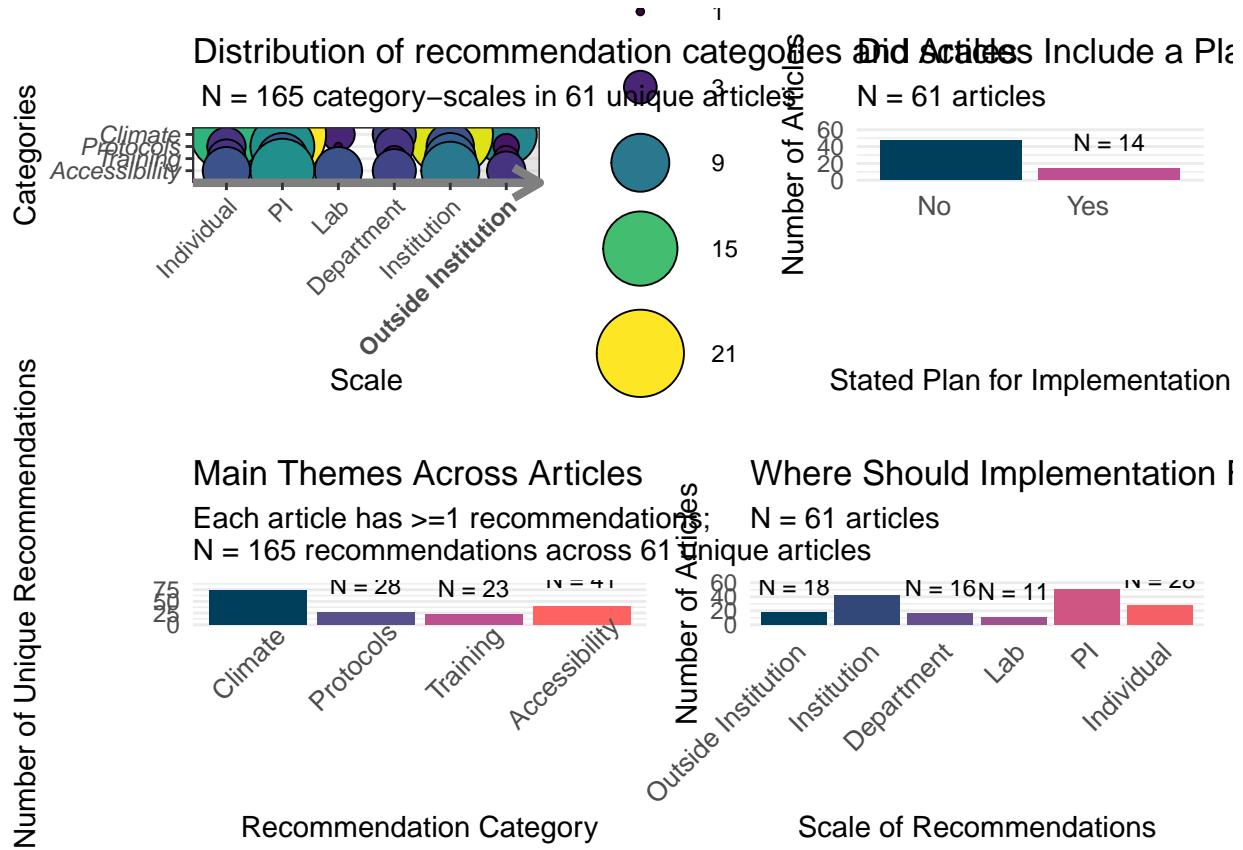
theme(legend.position = "none",
      panel.grid.major.x = element_blank(),
      axis.text.x = element_text(angle = 45, hjust = 1, size = 10),
      panel.background = element_rect(fill = "white", color = NA),
      plot.background = element_rect(fill = "white", color = NA))

# responsibility
# ggsave("Implementation Level.png", width = 6, height = 4)

```

COMBINED PLOTS

```
(heatmap + plan) / plot_spacer() / (themes + responsibility) +
  plot_layout(heights = c(3, 0.25, 2.5))
```



```
# ggsave("combined_plots.png", height = 10, width = 12)
```

Sankey

PLOT

```
pal6 <- as.list(pal(6))
pal6 <- unlist(pal6)

# Prepare alluvial data
alluvial_data <- q %>%
  select(
    category = recommendation_categories,
    scale = recommendation_scale,
    plan = plan_for_implementation_y_n
  ) %>%
  na.omit() %>%
  mutate(
    category = fct_relevel(category, c("Climate", "Protocols", "Training", "Accessibility")),
    scale = factor(scale, levels = c("Outside Institution", "Institution", "Department", "PI", "Lab", "N/A")),
    plan = factor(plan, levels = c("Y", "N"))
  ) %>%
  count(category, scale, plan, name = "n") %>%
  mutate(alluvium_id = paste(category, scale, sep = " → "))

# assign color palette to scales instead of categories
scale_levels <- levels(alluvial_data$scale)
scale_palette <- setNames(pal6[1:length(scale_levels)], scale_levels)

# plot with wider strata and custom colors
ggplot(alluvial_data,
       aes(axis1 = category, axis2 = scale, axis3 = plan, y = n)) +
  geom_flow(aes(fill = scale), width = 0.25, alpha = 0.75) +
  geom_stratum(width = 0.25, lwd = 0.4, color = "black") +
  geom_text(
    stat = "stratum",
    aes(
      label = after_stat(stratum),
      angle = dplyr::case_when(
        after_stat(x) %in% c(1, 2) ~ 20,
        TRUE ~ 0
      ),
      hjust = 0.5
    ),
    size = 2.5
  ) +
  scale_x_discrete(limits = c("Category", "Scale", "Plan"), expand = c(0.1, 0.1)) +
  scale_fill_manual(values = scale_palette) +
  labs(
    title = "Flow of Recommendations: Category → Scale → Implementation Plan",
    y = "Number of Mentions",
    x = NULL,
    fill = "Scale",
```

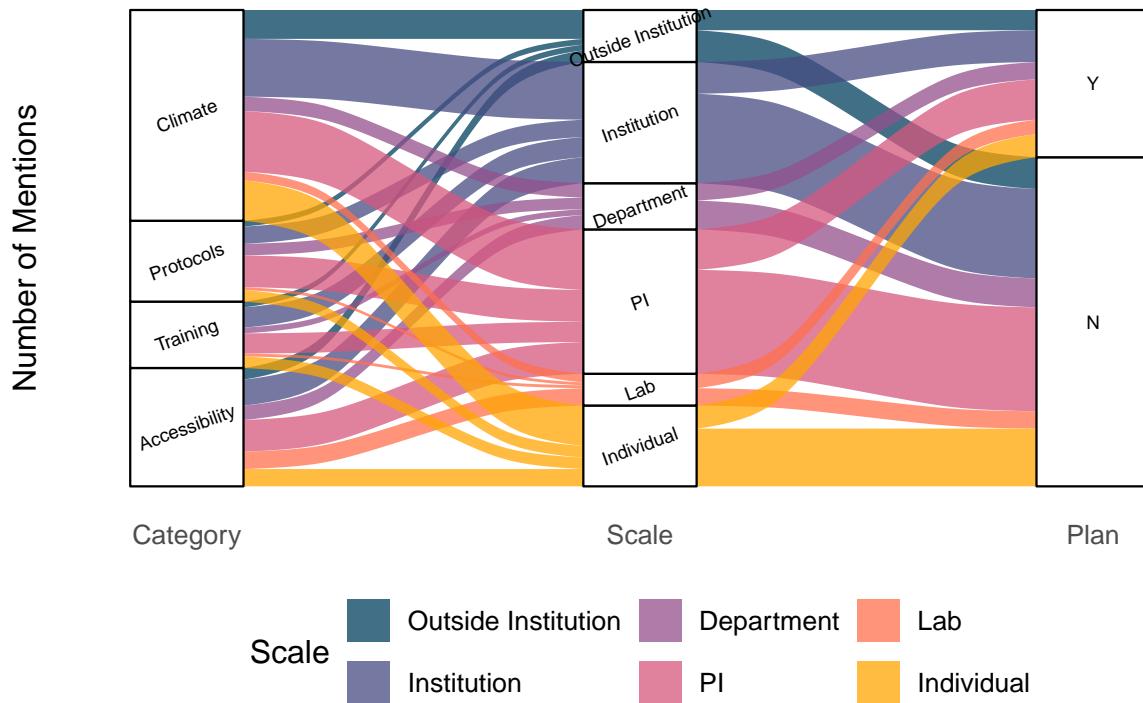
```

    subtitle = paste0("N = ", q_unique_recs_num, " unique recommendations across ", q_unique_articles,
) +
theme_minimal(base_size = 12) +
theme(
  legend.position = "bottom",
  plot.title = element_text(size = 14, face = "bold"),
  axis.text.y = element_blank(),
  axis.ticks.y = element_blank(),
  panel.grid = element_blank(),
  panel.background = element_rect(fill = "white", color = NA),
  plot.background = element_rect(fill = "white", color = NA)
) +
guides(fill = guide_legend(override.aes = list(size = 1)))

```

Flow of Recommendations: Category → Scale → Implementation P

N = 165 unique recommendations across 61 articles



```
# ggsave("flow_of_recs.png", width = 8, height = 6)
```

Disproportionality

The disproportionality analysis tests whether certain recommendation categories or scales were over- or under-represented among those with stated implementation plans.

```
### Calculate Skew - CATEGORY
```

```

# calculate overall baseline proportions of categories across all recommendations
category_proportions <- q %>%
  count(recommendation_categories, name = "total_n") %>%
  mutate(category_total_prop = total_n / sum(total_n)) %>%
  select(recommendation_categories, category_total_prop)

# calculate proportions within each plan group (Y/N)
category_plan_distribution <- q %>%
  mutate(plan = factor(plan_for_implementation_y_n, levels = c("Y", "N"))) %>%
  count(recommendation_categories, plan, name = "plan_n") %>%
  group_by(plan) %>%
  mutate(plan_total = sum(plan_n),
         category_within_plan_prop = plan_n / plan_total) %>%
  ungroup()

# join proportions and calculate difference (disproportionality)
comparison_by_category <- category_plan_distribution %>%
  left_join(category_proportions, by = "recommendation_categories") %>%
  mutate(disproportionality = category_within_plan_prop - category_total_prop) %>%
  arrange(recommendation_categories, plan)

# format as percentages
comparison_by_category_formatted <- comparison_by_category %>%
  mutate(across(
    c(category_total_prop, category_within_plan_prop, disproportionality),
    ~ percent(.x, accuracy = 0.1)
  ))

comparison_by_category_formatted

## # A tibble: 8 x 7
##   recommendation_categories plan  plan_n plan_total category_within_plan_prop
##   <fct>                  <fct> <int>     <int> <chr>
## 1 Climate                 Y      18       114 35.3%
## 2 Climate                 N      55       114 48.2%
## 3 Protocols                Y      14       51 27.5%
## 4 Protocols                N      14       114 12.3%
## 5 Training                 Y      8        51 15.7%
## 6 Training                 N      15       114 13.2%
## 7 Accessibility             Y      11       51 21.6%
## 8 Accessibility             N      30       114 26.3%
## # i 2 more variables: category_total_prop <chr>, disproportionality <chr>

write_csv(comparison_by_category_formatted, "comparison_ordered_category Nov 2025.csv")

```

Calculate Skew - SCALE

```

# factor order
scale_order <- c("Individual", "Lab", "PI", "Department", "Institution", "Outside Institution")

# calculate compute overall baseline proportions by scale

```

```

scale_proportions <- q %>%
  mutate(recommendation_scale = factor(recommendation_scale, levels = scale_order)) %>%
  count(recommendation_scale) %>%
  mutate(scale_total_prop = n / sum(n)) %>%
  select(recommendation_scale, scale_total_prop)

# calculate within-plan proportions by scale
scale_plan_distribution <- q %>%
  mutate(
    recommendation_scale = factor(recommendation_scale, levels = scale_order),
    plan = factor(plan_for_implementation_y_n, levels = c("Y", "N"))
  ) %>%
  count(recommendation_scale, plan) %>%
  group_by(plan) %>%
  mutate(plan_total = sum(n),
    scale_within_plan_prop = n / plan_total) %>%
  ungroup()

# join and calculate disproportionality between within-plan and overall proportions
comparison_ordered <- scale_plan_distribution %>%
  left_join(scale_proportions, by = "recommendation_scale") %>%
  mutate(disproportionality = scale_within_plan_prop - scale_total_prop) %>%
  arrange(recommendation_scale, plan) %>%
  mutate(across(
    c(scale_total_prop, scale_within_plan_prop, disproportionality),
    percent_format(accuracy = 0.1)
  ))

comparison_ordered

## # A tibble: 12 x 7
##   recommendation_scale plan      n plan_total scale_within_plan_prop
##   <fct>              <fct> <int>     <int> <chr>
## 1 Individual          Y        8       51  15.7%
## 2 Individual          N       20      114 17.5%
## 3 Lab                 Y        5       51  9.8%
## 4 Lab                 N        6      114 5.3%
## 5 PI                  Y       14      51 27.5%
## 6 PI                  N       36      114 31.6%
## 7 Department          Y        6       51 11.8%
## 8 Department          N       10      114 8.8%
## 9 Institution         Y       11      51 21.6%
## 10 Institution        N       31      114 27.2%
## 11 Outside Institution Y        7       51 13.7%
## 12 Outside Institution N       11      114 9.6%
## # i 2 more variables: scale_total_prop <chr>, disproportionality <chr>

write_csv(comparison_ordered, "comparison_ordered_scale Nov 2025.csv")

```