```
library(tidyverse)
## -- Attaching packages -----
1.3.0 --
## v ggplot2 3.3.3 v purrr 0.3.4
## v tibble 3.1.0 v dplyr 1.0.5
## v tidyr 1.1.3 v stringr 1.4.0
## v readr 1.4.0 v forcats 0.5.1
## -- Conflicts -----
                                      ----- tidyverse_conflicts()
## x \ dplyr::filter() \ masks \ stats::filter()
## x dplyr::lag() masks stats::lag()
library(targets)
library(glue)
##
## Attaching package: 'glue'
## The following object is masked from 'package:dplyr':
##
##
     collapse
withr::with_dir(here::here(), {
 tar_load(hpp_dat)
 tar_load(models)
})
```

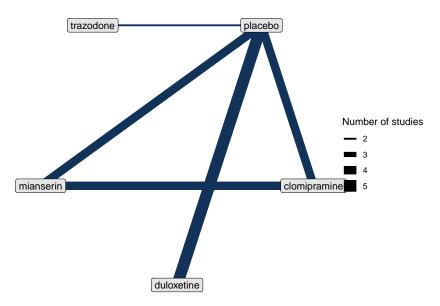
Hello, world.

#### 1 networks

```
models %>%
  map(.f = function(this_model) {
    outcome <- this_model %>% pluck("outcome")

  this_model %>%
    pluck("network") %>%
    plot() +
    labs(
        title = outcome,
        subtitle = "direct evidence comparisons"
    )
})
```

## substantial pain



```
## ## $model_binom_df381d93
```

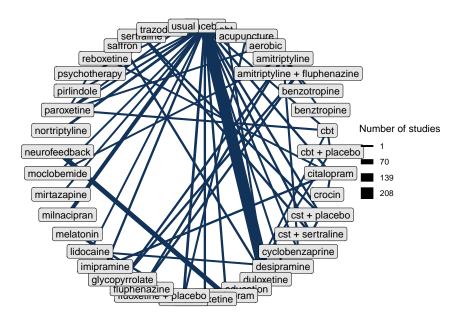
# moderate pain



```
##
## $model_smd_a56a0319
```

#### mood

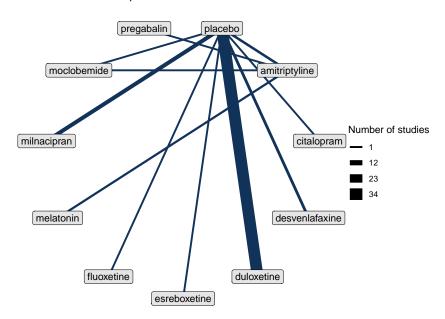
direct evidence comparisons



## ## \$model\_smd\_729faf21

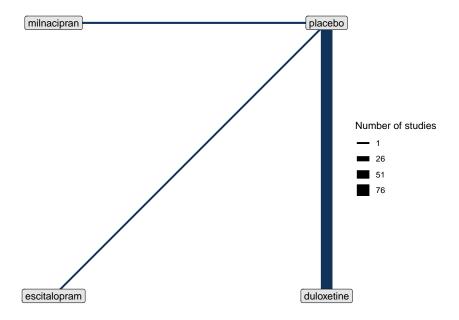
### sleep

direct evidence comparisons



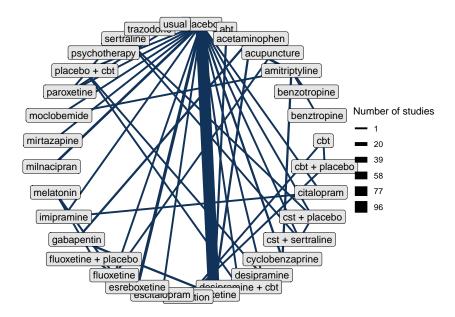
## ## \$model\_smd\_364aa3b7

pgic

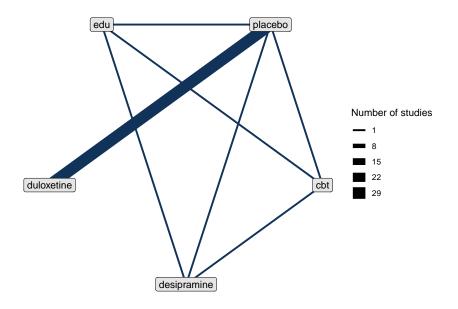


```
##
## $model_smd_cff505ca
```

### physical function



**qol** direct evidence comparisons



```
##
## $model_smd_1e5fe681
```

pain

