

Data Model SPZ

```
library(knitr)
library(readxl)
library(dplyr)
```

Attaching package: 'dplyr'

The following objects are masked from 'package:stats':

`filter`, `lag`

The following objects are masked from 'package:base':

`intersect`, `setdiff`, `setequal`, `union`

```
library(ggplot2)
library(gt)
library(treemap)
library(ggstatsplot)
```

You can cite this package as:

Patil, I. (2021). Visualizations with statistical details: The 'ggstatsplot' approach. Journal of Open Source Software, 6(61), 3167, doi:10.21105/joss.03167

```
library(grid) # for unit()
library(RColorBrewer)
library(ggsci)
library(ggpubr)
```

```
library(forcats)
library(icd.data)
library(tidyr)

colores<-get_palette(palette="lancet",12)
get_palette(palette="futurama",12)
```

```
[1] "#FF6F00FF" "#C71000FF" "#008EA0FF" "#8A4198FF" "#5A9599FF" "#FF6348FF"
[7] "#84D7E1FF" "#FF95A8FF" "#3D3B25FF" "#ADE2D0FF" "#1A5354FF" "#3F4041FF"
```

```
get_palette(palette="simpsons",12)
```

```
[1] "#FED439FF" "#709AE1FF" "#8A9197FF" "#D2AF81FF" "#FD7446FF" "#D5E4A2FF"
[7] "#197EC0FF" "#F05C3BFF" "#46732EFF" "#71D0F5FF" "#370335FF" "#075149FF"
```

```
get_palette(palette="tron",12)
```

```
[1] "#FF410D" "#AF9891" "#7ADFEC" "#C5CF7B" "#E5C638" "#AFCA51" "#A5D183"
[8] "#C5DBCD" "#DEC69D" "#F3A230" "#BB945B" "#748AA6"
```

```
#paleta<-get_palette(palette="simpsons",k = k)
```

```
#We upload data
codes <- read_excel("C:/Users/boris/Desktop/data hospital/codes_2phases.xlsx")
head(codes)
```

```
# A tibble: 6 x 3
  code1   date      code2
  <chr>   <dtm>      <chr>
1 6012487 2017-12-07 00:00:00 6014266
2 6023881 2018-12-06 00:00:00 6025255
3 6028398 2019-04-09 00:00:00 6028919
4 6040865 2020-08-10 00:00:00 6041525
5 6043203 2020-10-30 00:00:00 6043916
6 6017204 2018-04-30 00:00:00 6018809
```

```
# SCIM values 2017-2020
```

```
SCIM_17_20 <- read_excel("C:/Users/boris/Desktop/data hospital/Auswertung_SCIM_2017-20201.
```

Warning: Expecting logical in F4131 / R4131C6: got 'ja'

```
colnames(SCIM_17_20)
```

[1] "NR"	"PATIENT_NR"
[3] "FALL_NR"	"FID"
[5] "KURZZEICHEN"	"LASTMEASURE"
[7] "SP_NAHRUNG"	"SP_WASCHEN_OBER"
[9] "SP_WASCHEN_UNTER"	"SP_ANZIEHEN_OBER"
[11] "SP_ANZIEHEN_UNTER"	"SP_GESICHT"
[13] "SP_SUB"	"AS_ATMUNG"
[15] "AS_BLASE"	"AS_DARM"
[17] "AS_TOILETTE"	"AS_SUB"
[19] "MH_BETT"	"MH_TRANSFER_BETT"
[21] "MH_TRANSFER_WC"	"MH_SUB"
[23] "MA_HAUS"	"MA_MITTLERE_DISTANZ"
[25] "MA_AUSSER_HAUS"	"MA_TREPPEN"
[27] "MA_TRANSFER_AUTO"	"MA_TRANSFER_BODEN"
[29] "MA_SUB"	"SCIM_TOTAL"
[31] "DATUM"	"ERZEUGER"
[33] "ERZEUGUNGSDATUM"	"AENDERER"
[35] "AENDERUNGSDATUM"	"ERZ_CLIENT_INFO"
[37] "ERSTELLER_ERGO"	"ERSTELLER_PFLEGE"
[39] "ERSTELLER_PHYSIO"	"SP_NAHRUNG_AE"
[41] "SP_WASCHEN_OBER_AE"	"SP_WASCHEN_UNTER_AE"
[43] "SP_ANZIEHEN_OBER_AE"	"SP_ANZIEHEN_UNTER_AE"
[45] "SP_GESICHT_AE"	"AS_ATMUNG_AE"
[47] "AS_BLASE_AE"	"AS_DARM_AE"
[49] "AS_TOILETTE_AE"	"MH_BETT_AE"
[51] "MH_TRANSFER_BETT_AE"	"MH_TRANSFER_WC_AE"
[53] "MA_HAUS_AE"	"MA_MITTLERE_DISTANZ_AE"
[55] "MA_AUSSER_HAUS_AE"	"MA_TREPPEN_AE"
[57] "MA_TRANSFER_AUTO_AE"	"MA_TRANSFER_BODEN_AE"
[59] "KF_VERSTEHEN_AE"	"KF_VERSTEHEN"
[61] "KF_KOMMUNIKATION_VERSTEHEN"	"KF_KOMMUNIKATION_VERSTEHEN_AE"
[63] "KF_AUSDRUCK"	"KF_AUSDRUCK_AE"
[65] "KF_KOMMUNIKATION_AUSDRUCK"	"KF_KOMMUNIKATION_AUSDRUCK_AE"

```
[67] "KF_SOZIALES_VERHALTEN"      "KF_SOZIALES_VERHALTEN_AE"
[69] "KF_PROBLEM_LOESUNG"        "KF_PROBLEM_LOESUNG_AE"
[71] "KF_SUB"
```

```
# We convert the patient code to character
```

```
SCIM_17_20<-SCIM_17_20 %>% mutate(FID=as.character(FID))
head(SCIM_17_20)
```

```
# A tibble: 6 x 71
  NR PATIENT_NR FALL_NR FID KURZZ~1 LASTM~2 SP_NA~3 SP_WA~4 SP_WA~5 SP_AN~6
  <dbl>      <dbl>   <dbl> <chr> <chr>   <lg1>   <chr>   <chr>   <chr>   <chr>
1 36604      73510     34 5205~ CUS     NA      0      0      0      0
2 38222      396796     2 5207~ HON     NA      1      0      0      0
3 38801      398190     3 5208~ ROBA    NA      2      1      0      0
4 39684      399299     2 5211~ PRB     NA      0      0      0      0
5 39967      400212     1 5211~ JME     NA      0      0      0      0
6 39942      399420     3 5211~ KAJE    NA      2      0      0      0
# ... with 61 more variables: SP_ANZIEHEN_UNTER <chr>, SP_GESICHT <chr>,
# SP_SUB <chr>, AS_ATMUNG <chr>, AS_BLAUSE <chr>, AS_DARM <chr>,
# AS_TOILETTE <chr>, AS_SUB <chr>, MH_BETT <chr>, MH_TRANSFER_BETT <chr>,
# MH_TRANSFER_WC <chr>, MH_SUB <chr>, MA_HAUS <chr>,
# MA_MITTLERE_DISTANZ <chr>, MA_AUSSER_HAUS <chr>, MA_TREPPEN <chr>,
# MA_TRANSFER_AUTO <chr>, MA_TRANSFER_BODEN <chr>, MA_SUB <chr>,
# SCIM_TOTAL <chr>, DATUM <dtm>, ERZEUGER <chr>, ERZEUGUNGSDATUM <dtm>, ...
```

From the previous data set we need to know that the key variable that identifies every row is the column FID. Note that FID means case identification number.

- FID: Fallidentifikationsnummer

```
SCIM_17_20 %>% select(FID)%>%table() %>%data.frame()%>%select(Freq)%>% table()
```

```
Freq
  1    2
1119 3240
```

From the last outcome we observe that there are FID repeated at most twice. It probably means two different SCIM values. In fact Lets select the first five FID values.

```
fives<-SCIM_17_20 %>% select(FID)%>%table() %>%data.frame()%>% head(n=5) %>% select(FID)
```

```
fives
```

```
FID
```

```
1 5205639
2 5207178
3 5208756
4 5210382
5 5211089
```

```
SCIM_17_20 %>% filter(FID %in% fives$FID)%>%select(FID,SCIM_TOTAL,DATUM) %>% arrange(FID,D
```

```
# A tibble: 10 x 3
```

	FID	SCIM_TOTAL	DATUM
	<chr>	<chr>	<dtm>
1	5205639	15	2015-10-29 00:00:00
2	5205639	21	2017-08-16 00:00:00
3	5207178	17	2016-01-01 00:00:00
4	5207178	42	2017-04-25 00:00:00
5	5208756	14	2016-03-01 00:00:00
6	5208756	63	2017-02-08 00:00:00
7	5210382	14	2017-04-11 00:00:00
8	5210382	14	2017-04-19 00:00:00
9	5211089	12	2016-05-01 00:00:00
10	5211089	84	2017-03-01 00:00:00

Analysis two phases

Here we consider the dataset *codes*, remember that the data set *codes* has two columns *code1* and *code2*. Both codes corresponds to the same **patient**, in different phases. Therefore, is important to keep in mind this feature because based on these codes we will analyze the evolution of SCIM values.

- code 1: Acute Phase
- code 2: Rehabilitation Phase

```
head(codes)
```

```
# A tibble: 6 x 3
  code1   date                code2
  <chr>   <dtm>                  <chr>
1 6012487 2017-12-07 00:00:00 6014266
2 6023881 2018-12-06 00:00:00 6025255
3 6028398 2019-04-09 00:00:00 6028919
4 6040865 2020-08-10 00:00:00 6041525
5 6043203 2020-10-30 00:00:00 6043916
6 6017204 2018-04-30 00:00:00 6018809
```

Additionally, is crucial to note that:

- There are not duplicated values in the table *codes*
- There are duplicated FID in the table *SCIM_17_20*.

Information about patients, diagnosis and treatments.

In this stage we consider two additional data sets.

The **mb** is the dataset with information about health diagnosis, treatments, age, sex among others.

The data set **fid_pid** is a data set that contains a correspondence between FID and PID.

- PID: Patienten fallnummer that is the patient case number.

```
mb<-readRDS("C:/Users/boris/Desktop/SPZ_hospital/SPZ-Project/mb.rds")
fid_pid <- readRDS("C:/Users/boris/Desktop/SPZ_hospital/SPZ-Project/fidpid.rds")
head(mb)
```

	les_lev	gender	birth_date	age	residence	stay_ICU	entry_date	leave_date
1	G8260	1	19630105	55	9462	0	2018091300	2018100823
2	G8260	1	19690920	50	3018	0	2019122007	2020010923
3	G8260	1	19550810	64	8800	0	2020011107	2020012511
4	G8260	2	19870614	32	8630	0	2019121800	2020022110
5	G8260	1	19800523	39	4492	0	2019092307	2019092711
6	G8260	1	19671106	51	2000	0	2019022007	2019022116
	cost_payer	main_diag	add_diag	sec_diag1	sec_diag2	sec_diag3	sec_diag4	
1	1	G950	<NA>	S4221	G8242	G8260	M8128	
2	1	G8230	<NA>	G8260	Z488	M4642	G9582	
3	4	G8242	<NA>	G8260	Z508	N210	N3188	

4	1	L8925	<NA>	B956	G8243	G8260	N210
5	4	K642	<NA>	G8242	G8260	Z881	I1090
6	4	G4738	<NA>	Z991	G8243	G8260	Z435
	sec_diag5	sec_diag6	sec_diag7	sec_diag8	main_treat	start_main_treat	add_main
1	Z223	B962	<NA>	<NA>	371	2018100807	99C110
2	J9609	Z430	A490	B956	93872	2019122007	93399
3	J9610	T835	Y828	B965	93872	2020011107	9319
4	G9584	Z435	<NA>	<NA>	862A1E	2019122707	99C12G
5	<NA>	<NA>	<NA>	<NA>	49461	2019092307	99C11
6	G9584	Z881	<NA>	<NA>	89381	2019022007	89381
	fur_treat1	fur_treat2	fur_treat3	fur_treat4	fur_treat5	fur_treat6	fur_treat7
1	889410	883820	889410	889714	889810	883840	30926
2	9383	890A3	93701	311	4311	89381	<NA>
3	93399	9383	570X1	5994	AA211	AA321	AA322
4	867E1E	570X10	5994	AA2117	AA3211	AA3221	AA3231
5	009A1	991	<NA>	<NA>	<NA>	<NA>	<NA>
6	8922	5732	<NA>	<NA>	<NA>	<NA>	<NA>
	fur_treat8	id	year	ind_level	Lesion	length_stay	
1	36	6022710	2018	1	C1-C3	25 days	
2	<NA>	6035865	2019	1	C1-C3	20 days	
3	AA323	6035854	2019	1	C1-C3	14 days	
4	AA3241	6035716	2020	1	C1-C3	65 days	
5	<NA>	6032762	2019	1	C1-C3	4 days	
6	<NA>	6026011	2019	1	C1-C3	1 days	

```
fid_pid%>%select(PID)%>%table()%>%data.frame()%>%select(Freq)%>% table()|> data.frame()
```

	Freq	Freq.1
1	1	1664
2	2	888
3	3	258
4	4	125
5	5	76
6	6	45
7	7	21
8	8	11
9	9	10
10	10	5
11	11	3
12	12	1
13	13	1
14	16	1

```
colnames(mb)
```

```
[1] "les_lev"          "gender"          "birth_date"      "age"
[5] "residence"        "stay_ICU"        "entry_date"      "leave_date"
[9] "cost_payer"       "main_diag"       "add_diag"        "sec_diag1"
[13] "sec_diag2"        "sec_diag3"       "sec_diag4"       "sec_diag5"
[17] "sec_diag6"        "sec_diag7"       "sec_diag8"       "main_treat"
[21] "start_main_treat" "add_main"        "fur_treat1"      "fur_treat2"
[25] "fur_treat3"       "fur_treat4"      "fur_treat5"      "fur_treat6"
[29] "fur_treat7"       "fur_treat8"      "id"              "year"
[33] "ind_level"        "Lesion"          "length_stay"
```

Variables time and cost of interventions

We include variables that represent the time and cost for each intervention to every patient

```
time_cost<-readRDS("time_cost.rds")
serv_cons<-time_cost%>% select(-year) %>% group_by(id) %>%
  summarize(op_room=sum(op_room),anaest=sum(anaest),int_care=sum(int_care),imag=sum(imag),
```

The units are given by:

```
unts<-c("min","min","min","chf","chf","min","min","min","tp","min","tp")
unts
```

```
[1] "min" "min" "min" "chf" "chf" "min" "min" "min" "tp" "min" "tp"
```

Summarizing, we have the next important data sets.

- mb
- codes
- SCIM_17_20
- fid_pid
- time_cost

Evolution of SCIM

```
SCIM_17_20 %>% select(FID) %>% table() %>% data.frame()%>% arrange(-Freq)%>%head(n=5)
```

	FID	Freq
1	5205639	2
2	5207178	2
3	5208756	2
4	5210382	2
5	5211089	2

```
mb%>% select(id) %>% table() %>% data.frame()%>%arrange(-Freq) %>% head(n=5)
```

	id	Freq
1	6006658	2
2	6008272	2
3	6008830	2
4	6009414	2
5	6009503	2

```
time_cost %>% select(id) %>% table() %>% data.frame() %>% arrange(-Freq) %>% head(n=8)
```

	id	Freq
1	5211529	2
2	6006658	2
3	6008272	2
4	6008830	2
5	6009414	2
6	6009503	2
7	6009636	2
8	6009808	2

```
fid_pid %>% select(FID) %>% table() %>% data.frame()%>% arrange(-Freq) %>% head(n=8)
```

	FID	Freq
1	5205639	1
2	5207178	1

```
3 5208756    1
4 5210382    1
5 5211089    1
6 5211529    1
7 5211936    1
8 5212187    1
```

```
fid_pid %>% select(PID) %>% table() %>% data.frame()%>% arrange(-Freq) %>% head(n=8)
```

```
      PID Freq
1 0152021   16
2 0019699   13
3 0001323   12
4 0000374   11
5 0002657   11
6 0153996   11
7 0001984   10
8 0002447   10
```

From the previous summaries we observe that in the table *fid_pid*, there are some *PID* codes repeated 16 times. It means that the patient considered was \$16\$ times in the clinic, each time with a different FID code.

```
fid_pid |> filter(PID=="0152021")
```

```
# A tibble: 16 x 2
  FID      PID
  <chr>   <chr>
1 5211529 0152021
2 6009261 0152021
3 6011127 0152021
4 6012509 0152021
5 6013263 0152021
6 6015622 0152021
7 6021794 0152021
8 6023916 0152021
9 6025811 0152021
10 6027140 0152021
11 6029079 0152021
12 6031905 0152021
```

```

13 6034556 0152021
14 6035696 0152021
15 6037398 0152021
16 6040704 0152021

```

```
fid_pid |> filter(PID=="0152021") |> select(FID) |> unique() |> nrow()
```

```
[1] 16
```

In order to analyze the evolution of the SCIM value we consider the next steps.

Due to the importance of track the acute and rehabilitation phase we merge the mb dataset with codes. Note that an important issue could be the treatment of the codes for acute and rehabilitation phase. In order to deal with that we proceed as follows:

```

# We create a code uni_n in the codes table
codes$uniq<-paste("uni",1:nrow(codes),sep="_")
head(codes,n=10)

```

```

# A tibble: 10 x 4
   code1   date                code2   uniq
   <chr>   <dtm>                <chr>   <chr>
1 6012487 2017-12-07 00:00:00 6014266 uni_1
2 6023881 2018-12-06 00:00:00 6025255 uni_2
3 6028398 2019-04-09 00:00:00 6028919 uni_3
4 6040865 2020-08-10 00:00:00 6041525 uni_4
5 6043203 2020-10-30 00:00:00 6043916 uni_5
6 6017204 2018-04-30 00:00:00 6018809 uni_6
7 6040437 2020-07-27 00:00:00 6041208 uni_7
8 6016770 2018-03-28 00:00:00 6017875 uni_8
9 6013568 2017-12-18 00:00:00 6014569 uni_9
10 6025398 2019-01-29 00:00:00 6026686 uni_10

```

```
SCIM_17_20 %>% select(FID) %>% unique() %>% nrow()
```

```
[1] 4359
```

```
# Note that when we merge SCIM_17_20 with codes not all codes match, because
# not all of them were in two phases
```

```
SCIM_17_20 %>% select(FID) %>%
  merge(codes%>%mutate(ph="acute"),by.x="FID",by.y="code1") %>% nrow()
```

[1] 1638

```
#1638 matches in acute phase
```

```
# Merge with code 1 acute phase
P1 <- SCIM_17_20 %>%
  merge(codes,by.x="FID",by.y="code1",all.x=TRUE)
```

```
head(P1)
```

	FID	NR	PATIENT_NR	FALL_NR	KURZZEICHEN	LASTMEASURE	SP_NAHRUNG	
1	5205639	36604	73510	34	CUS	NA	0	
2	5205639	46713	73510	34	PUNO	NA	0	
3	5207178	38222	396796	2	HON	NA	1	
4	5207178	44542	396796	2	SILG	NA	2	
5	5208756	38801	398190	3	ROBA	NA	2	
6	5208756	42935	398190	3	ENBE	NA	3	
	SP_WASCHEN_OBER	SP_WASCHEN_UNTER	SP_ANZIEHEN_OBER	SP_ANZIEHEN_UNTER				
1	0		0		0		0	
2	0		0		0		0	
3	0		0		0		0	
4	0		0		1		1	
5	1		0		0		0	
6	3		1		4		2	
	SP_GESICHT	SP_SUB	AS_ATMUNG	AS_BLASE	AS_DARM	AS_TOILETTE	AS_SUB	MH_BETT
1	0	0	10	0	5	0	15	0
2	0	0	10	3	5	0	18	0
3	1	2	4	3	5	0	12	0
4	0	4	10	6	10	1	27	2
5	1	4	10	0	0	0	10	0
6	3	16	10	11	10	5	36	4
	MH_TRANSFER_BETT	MH_TRANSFER_WC	MH_SUB	MA_HAUS	MA_MITTLERE_DISTANZ			
1	0		0	0			0	
2	0		0	0	1			1

3	0	0	0	1	1
4	2	1	5	2	2
5	0	0	0	0	0
6	2	1	7	1	1
MA_AUSSER_HAUS MA_TREPPEN MA_TRANSFER_AUTO MA_TRANSFER_BODEN MA_SUB					
1	0	0	0	0	0
2	1	0	0	0	3
3	1	0	0	0	3
4	1	0	1	0	6
5	0	0	0	0	0
6	1	0	1	0	4
SCIM_TOTAL DATUM ERZEUGER ERZEUGUNGSDATUM AENDERER AENDERUNGSDATUM					
1	15	2015-10-29	SMITH_C	2015-10-29 22:15:23	NA NA
2	21	2017-08-16	PUERRO_N	2017-08-16 08:10:28	NA NA
3	17	2016-01-01	HOFER_N	2015-12-31 07:57:54	NA NA
4	42	2017-04-25	GRETHER_S	2017-04-25 14:59:53	NA NA
5	14	2016-03-01	BACHMANN_R	2016-02-26 11:13:36	NA NA
6	63	2017-02-08	ENDERS_B	2017-02-06 08:37:41	NA NA
ERZ_CLIENT_INFO ERSTELLER_ERGO ERSTELLER_PFLEGE					
1	KgEd32@SPGVCMP00026-3.0.0.0-2.2.0.1921			JUST	CUS
2	KgEd32@SPG16010-3.0.0.0-2.2.0.2143			CLE	PUNO
3	KgEd32@SPGVCMP00040-3.0.0.0-2.2.0.2027			ANWA	REC
4	KgEd32@SPG01806-3.0.0.0-2.2.0.2143			SILG	NIV
5	KgEd32@SPG01734-3.0.0.0-2.2.0.2054			FENI	BYZY
6	KgEd32@SPGPVCMP00065-3.0.0.0-2.2.0.2085			ENBE	BER
ERSTELLER_PHYSIO SP_NAHRUNG_AE SP_WASCHEN_OBER_AE SP_WASCHEN_UNTER_AE					
1	KOST	CUS	CUS	CUS	
2	KOST	PUNO	PUNO	PUNO	
3	HON	REC	REC	REC	
4	SILG	NIV	NIV	NIV	
5	ROSE	BYZY	BYZY	BYZY	
6	JED	BER	BER	BER	
SP_ANZIEHEN_OBER_AE SP_ANZIEHEN_UNTER_AE SP_GESICHT_AE AS_ATMUNG_AE					
1	CUS	CUS	CUS	CUS	
2	PUNO	PUNO	PUNO	PUNO	
3	REC	REC	REC	REC	
4	NIV	NIV	NIV	NIV	
5	BYZY	BYZY	BYZY	BYZY	
6	BER	BER	BER	BER	
AS_BLAASE_AE AS_DARM_AE AS_TOILETTE_AE MH_BETT_AE MH_TRANSFER_BETT_AE					
1	CUS	CUS	CUS	CUS	JUST
2	PUNO	PUNO	PUNO	PUNO	CLE
3	REC	REC	REC	REC	ANWA

4	NIV	NIV	NIV	NIV	SILG
5	BYZY	BYZY	BYZY	BYZY	FENI
6	BER	BER	BER	BER	ENBE
	MH_TRANSFER_WC_AE	MA_HAUS_AE	MA_MITTLERE_DISTANZ_AE	MA_AUSSER_HAUS_AE	
1	JUST	KOST		KOST	KOST
2	CLE	KOST		KOST	KOST
3	ANWA	HON		HON	HON
4	SILG	SILG		SILG	SILG
5	FENI	ROSE		ROSE	ROSE
6	ENBE	JED		JED	JED
	MA_TREPPEN_AE	MA_TRANSFER_AUTO_AE	MA_TRANSFER_BODEN_AE	KF_VERSTEHEN_AE	
1	KOST	JUST	KOST	<NA>	
2	KOST	CLE	KOST	CLE	
3	HON	ANWA	HON	<NA>	
4	SILG	SILG	SILG	<NA>	
5	ROSE	FENI	ROSE	<NA>	
6	JED	ENBE	JED	<NA>	
	KF_VERSTEHEN	KF_KOMMUNIKATION_VERSTEHEN	KF_KOMMUNIKATION_VERSTEHEN_AE		
1	<NA>		<NA>	<NA>	
2	7	3 = beides		CLE	
3	<NA>		<NA>	<NA>	
4	<NA>		<NA>	<NA>	
5	<NA>		<NA>	<NA>	
6	<NA>		<NA>	<NA>	
	KF_AUSDRUCK	KF_AUSDRUCK_AE	KF_KOMMUNIKATION_AUSDRUCK		
1	<NA>	<NA>	<NA>		
2	7	CLE	3 = beides		
3	<NA>	<NA>	<NA>		
4	<NA>	<NA>	<NA>		
5	<NA>	<NA>	<NA>		
6	<NA>	<NA>	<NA>		
	KF_KOMMUNIKATION_AUSDRUCK_AE	KF_SOZIALES_VERHALTEN	KF_SOZIALES_VERHALTEN_AE		
1		<NA>	<NA>	<NA>	
2		CLE	7	CLE	
3		<NA>	<NA>	<NA>	
4		<NA>	<NA>	<NA>	
5		<NA>	<NA>	<NA>	
6		<NA>	<NA>	<NA>	
	KF_PROBLEM_LOESUNG	KF_PROBLEM_LOESUNG_AE	KF_SUB	date	code2
1	<NA>	<NA>	<NA>	<NA>	<NA>
2	7	CLE	34	<NA>	<NA>
3	<NA>	<NA>	<NA>	<NA>	<NA>
4	<NA>	<NA>	-	<NA>	<NA>

```

5          <NA>          <NA>  <NA> <NA> <NA> <NA>
6          <NA>          <NA>    - <NA> <NA> <NA>

```

```
# Merge with code 2 rehabilitation phase
```

```
P2 <- P1 %>% merge(codes,by.x="FID",by.y="code2",all.x=TRUE)
```

```
# For example
```

```
head(P2[c(71,72,73,74,449,450,451,452),])
```

	FID	NR	PATIENT_NR	FALL_NR	KURZZEICHEN	LASTMEASURE	SP_NAHRUNG	
71	5215438	42899	9567	52	BUIR	NA	2	
72	5215438	42730	9567	52	ALLU	NA	2	
73	5215452	41365	393174	3	MAME	NA	3	
74	5215452	42776	393174	3	MAME	NA	3	
449	6002536	42900	170836	6	MEOS	NA	2	
450	6002536	47031	170836	6	JEDI	NA	2	
	SP_WASCHEN_OBER	SP_WASCHEN_UNTER	SP_ANZIEHEN_OBER	SP_ANZIEHEN_UNTER				
71	1	0	0	0				
72	0	0	0	0				
73	3	3	4	4				
74	3	3	4	4				
449	1	0	1	0				
450	3	2	2	1				
	SP_GESICHT	SP_SUB	AS_ATMUNG	AS_BLASE	AS_DARM	AS_TOILETTE	AS_SUB	MH_BETT
71	3	6	10	3	5	0	18	0
72	2	4	10	0	5	0	15	0
73	3	20	10	0	0	5	15	6
74	3	20	10	11	10	4	35	6
449	1	5	10	0	5	0	15	2
450	3	13	10	15	10	4	39	6
	MH_TRANSFER_BETT	MH_TRANSFER_WC	MH_SUB	MA_HAUS	MA_MITTLERE_DISTANZ			
71	0	0	0	0				
72	0	0	0	2				
73	2	1	9	2				
74	2	1	9	2				
449	0	0	2	0				
450	2	1	9	4				
	MA_AUSSER_HAUS	MA_TREPPEN	MA_TRANSFER_AUTO	MA_TRANSFER_BODEN	MA_SUB			
71	0	0	0	0	0			
72	2	0	0	0	0			
73	2	0	2	0	0			

74	2	0	2	1	9
449	0	0	0	0	0
450	4	2	1	0	15
	SCIM_TOTAL	DATUM	ERZEUGER	ERZEUGUNGSDATUM	AENDERER
71	24	2017-02-01	SCHAUB_I	2017-02-01 11:17:38	NA
72	25	2017-01-18	LUTZ_A	2017-01-20 14:25:42	NA
73	52	2016-09-19	MERLONI_M	2016-09-22 13:09:13	NA
74	73	2017-01-27	MERLONI_M	2017-01-27 16:47:49	NA
449	22	2017-02-01	OSELE_M	2017-02-01 12:58:12	NA
450	76	2017-08-28	DIENEMANN_J	2017-08-25 08:56:03	NA
	AENDERUNGSDATUM		ERZ_CLIENT_INFO	ERSTELLER_ERGO	
71	NA	KgEd32@SPGPVCMP00050-3.0.0.0-2.2.0.2085		ALLU	
72	NA	KgEd32@SPGPVCMP00035-3.0.0.0-2.2.0.2085		ALLU	
73	NA	KgEd32@SPGPVCMP00060-3.0.0.0-2.2.0.2085		MAME	
74	NA	KgEd32@SPGPVCMP00016-3.0.0.0-2.2.0.2085		MAME	
449	NA	KgEd32@SPGPVCMP00037-3.0.0.0-2.2.0.2085		MAME	
450	NA	KgEd32@SPGPVCMP00035-3.0.0.0-2.2.0.2143		MAME	
	ERSTELLER_PFLEGE	ERSTELLER_PHYSIO	SP_NAHRUNG_AE	SP_WASCHEN_OBER_AE	
71	LAC	BUIR	LAC	LAC	
72	LAU	BUIR	LAU	LAU	
73	BAE	LOSA	BAE	BAE	
74	BEIS	LOSA	BEIS	BEIS	
449	THAN	MEOS	THAN	THAN	
450	THAN	JEDI	THAN	THAN	
	SP_WASCHEN_UNTER_AE	SP_ANZIEHEN_OBER_AE	SP_ANZIEHEN_UNTER_AE	SP_GESICHT_AE	
71	LAC		LAC	LAC	LAC
72	LAU		LAU	LAU	LAU
73	BAE		BAE	BAE	BAE
74	BEIS		BEIS	BEIS	BEIS
449	THAN		THAN	THAN	THAN
450	THAN		THAN	THAN	THAN
	AS_ATMUNG_AE	AS_BLAASE_AE	AS_DARM_AE	AS_TOILETTE_AE	MH_BETT_AE
71	LAC	LAC	LAC	LAC	LAC
72	LAU	LAU	LAU	LAU	LAU
73	BAE	BAE	BAE	BAE	BAE
74	BEIS	BEIS	BEIS	BEIS	BEIS
449	THAN	THAN	THAN	THAN	THAN
450	THAN	THAN	THAN	THAN	THAN
	MH_TRANSFER_BETT_AE	MH_TRANSFER_WC_AE	MA_HAUS_AE	MA_MITTLERE_DISTANZ_AE	
71	ALLU		ALLU	BUIR	BUIR
72	ALLU		ALLU	BUIR	BUIR
73	MAME		MAME	LOSA	LOSA
74	MAME		MAME	LOSA	LOSA


```

450          MAME      34      <NA>      <NA>      <NA> 5217254 2017-01-11
      uniq.y
71      <NA>
72      <NA>
73      <NA>
74      <NA>
449 uni_247
450 uni_247

```

If everything is ok, when uniq.x has any value uniq.y should be with <NA> and viceversa. The next chunk evaluates if both columns uniq.x and uniq.y are not empty at the same time.

```

#P2 %>% filter(uniq.x!=""&uniq.y!="")
## 0 rows has both values empty at the same time

P2$uniq.x<-ifelse(is.na(P2$uniq.x),"",P2$uniq.x)
P2$uniq.y<-ifelse(is.na(P2$uniq.y),"",P2$uniq.y)

P2$unique<-paste(P2$uniq.x,P2$uniq.y,sep="")
head(P2)

```

	FID	NR	PATIENT_NR	FALL_NR	KURZZEICHEN	LASTMEASURE	SP_NAHRUNG	
1	5205639	36604	73510	34	CUS	NA	0	
2	5205639	46713	73510	34	PUNO	NA	0	
3	5207178	38222	396796	2	HON	NA	1	
4	5207178	44542	396796	2	SILG	NA	2	
5	5208756	38801	398190	3	ROBA	NA	2	
6	5208756	42935	398190	3	ENBE	NA	3	
	SP_WASCHEN_OBER	SP_WASCHEN_UNTER	SP_ANZIEHEN_OBER	SP_ANZIEHEN_UNTER				
1	0	0	0	0				
2	0	0	0	0				
3	0	0	0	0				
4	0	0	1	1				
5	1	0	0	0				
6	3	1	4	2				
	SP_GESICHT	SP_SUB	AS_ATMUNG	AS_BLASE	AS_DARM	AS_TOILETTE	AS_SUB	MH_BETT
1	0	0	10	0	5	0	15	0
2	0	0	10	3	5	0	18	0
3	1	2	4	3	5	0	12	0
4	0	4	10	6	10	1	27	2
5	1	4	10	0	0	0	10	0
6	3	16	10	11	10	5	36	4

	MH_TRANSFER_BETT	MH_TRANSFER_WC	MH_SUB	MA_HAUS	MA_MITTLERE_DISTANZ
1	0	0	0	0	0
2	0	0	0	1	1
3	0	0	0	1	1
4	2	1	5	2	2
5	0	0	0	0	0
6	2	1	7	1	1

	MA_AUSSER_HAUS	MA_TREPPEN	MA_TRANSFER_AUTO	MA_TRANSFER_BODEN	MA_SUB
1	0	0	0	0	0
2	1	0	0	0	3
3	1	0	0	0	3
4	1	0	1	0	6
5	0	0	0	0	0
6	1	0	1	0	4

	SCIM_TOTAL	DATUM	ERZEUGER	ERZEUGUNGSDATUM	AENDERER	AENDERUNGSDATUM
1	15	2015-10-29	SMITH_C	2015-10-29 22:15:23	NA	NA
2	21	2017-08-16	PUERRO_N	2017-08-16 08:10:28	NA	NA
3	17	2016-01-01	HOFER_N	2015-12-31 07:57:54	NA	NA
4	42	2017-04-25	GRETHER_S	2017-04-25 14:59:53	NA	NA
5	14	2016-03-01	BACHMANN_R	2016-02-26 11:13:36	NA	NA
6	63	2017-02-08	ENDERS_B	2017-02-06 08:37:41	NA	NA

	ERZ_CLIENT_INFO	ERSTELLER_ERGO	ERSTELLER_PFLEGE
1	KgEd32@SPGVCMP00026-3.0.0.0-2.2.0.1921	JUST	CUS
2	KgEd32@SPG16010-3.0.0.0-2.2.0.2143	CLE	PUNO
3	KgEd32@SPGVCMP00040-3.0.0.0-2.2.0.2027	ANWA	REC
4	KgEd32@SPG01806-3.0.0.0-2.2.0.2143	SILG	NIV
5	KgEd32@SPG01734-3.0.0.0-2.2.0.2054	FENI	BYZY
6	KgEd32@SPGPVCM00065-3.0.0.0-2.2.0.2085	ENBE	BER

	ERSTELLER_PHYSIO	SP_NAHRUNG_AE	SP_WASCHEN_OBER_AE	SP_WASCHEN_UNTER_AE
1	KOST	CUS	CUS	CUS
2	KOST	PUNO	PUNO	PUNO
3	HON	REC	REC	REC
4	SILG	NIV	NIV	NIV
5	ROSE	BYZY	BYZY	BYZY
6	JED	BER	BER	BER

	SP_ANZIEHEN_OBER_AE	SP_ANZIEHEN_UNTER_AE	SP_GESICHT_AE	AS_ATMUNG_AE
1	CUS	CUS	CUS	CUS
2	PUNO	PUNO	PUNO	PUNO
3	REC	REC	REC	REC
4	NIV	NIV	NIV	NIV
5	BYZY	BYZY	BYZY	BYZY
6	BER	BER	BER	BER

	AS_BLASE_AE	AS_DARM_AE	AS_TOILETTE_AE	MH_BETT_AE	MH_TRANSFER_BETT_AE
--	-------------	------------	----------------	------------	---------------------

1	CUS	CUS	CUS	CUS	JUST
2	PUNO	PUNO	PUNO	PUNO	CLE
3	REC	REC	REC	REC	ANWA
4	NIV	NIV	NIV	NIV	SILG
5	BYZY	BYZY	BYZY	BYZY	FENI
6	BER	BER	BER	BER	ENBE
	MH_TRANSFER_WC_AE	MA_HAUS_AE	MA_MITTLERE_DISTANZ_AE	MA_AUSSER_HAUS_AE	
1		JUST	KOST	KOST	KOST
2		CLE	KOST	KOST	KOST
3		ANWA	HON	HON	HON
4		SILG	SILG	SILG	SILG
5		FENI	ROSE	ROSE	ROSE
6		ENBE	JED	JED	JED
	MA_TREPPEN_AE	MA_TRANSFER_AUTO_AE	MA_TRANSFER_BODEN_AE	KF_VERSTEHEN_AE	
1		KOST	JUST	KOST	<NA>
2		KOST	CLE	KOST	CLE
3		HON	ANWA	HON	<NA>
4		SILG	SILG	SILG	<NA>
5		ROSE	FENI	ROSE	<NA>
6		JED	ENBE	JED	<NA>
	KF_VERSTEHEN	KF_KOMMUNIKATION_VERSTEHEN	KF_KOMMUNIKATION_VERSTEHEN_AE		
1		<NA>	<NA>		<NA>
2		7	3 = beides		CLE
3		<NA>	<NA>		<NA>
4		<NA>	<NA>		<NA>
5		<NA>	<NA>		<NA>
6		<NA>	<NA>		<NA>
	KF_AUSDRUCK	KF_AUSDRUCK_AE	KF_KOMMUNIKATION_AUSDRUCK		
1		<NA>	<NA>		<NA>
2		7	CLE	3 = beides	
3		<NA>	<NA>		<NA>
4		<NA>	<NA>		<NA>
5		<NA>	<NA>		<NA>
6		<NA>	<NA>		<NA>
	KF_KOMMUNIKATION_AUSDRUCK_AE	KF_SOZIALES_VERHALTEN	KF_SOZIALES_VERHALTEN_AE		
1		<NA>	<NA>		<NA>
2		CLE	7		CLE
3		<NA>	<NA>		<NA>
4		<NA>	<NA>		<NA>
5		<NA>	<NA>		<NA>
6		<NA>	<NA>		<NA>
	KF_PROBLEM_LOESUNG	KF_PROBLEM_LOESUNG_AE	KF_SUB	date.x	code2
1		<NA>	<NA>	<NA>	<NA>

2	7	CLE	34	<NA>	<NA>	<NA>
3	<NA>	<NA>	<NA>	<NA>	<NA>	<NA>
4	<NA>	<NA>	-	<NA>	<NA>	<NA>
5	<NA>	<NA>	<NA>	<NA>	<NA>	<NA>
6	<NA>	<NA>	-	<NA>	<NA>	<NA>

	date.y	uniq.y	unique
1	<NA>		
2	<NA>		
3	<NA>		
4	<NA>		
5	<NA>		
6	<NA>		

```
P2$acute<-ifelse(P2$uniq.x=="", "", "acute")
P2$rehab<-ifelse(P2$uniq.y=="", "", "rehab")
```

```
P2$phase<-paste(P2$acute,P2$rehab,sep="")
```

```
P3<-P2%>% select(unique,FID,code1,code2,phase,SCIM_TOTAL)
#head(P2)
#head(P3)
tabl<-table(P3$unique)%>% data.frame() %>% arrange(-Freq)
head(tabl)
```

	Var1	Freq
1		3976
2	uni_100	4
3	uni_1000	4
4	uni_1001	4
5	uni_1002	4
6	uni_1003	4

```
#P3%>%filter(unique %in% tabl[-1,]$Var1) %>% merge(fid_pid,by.x="FID",by.y="FID")%>% arrange(PID)
P4<-P3%>%filter(!is.na(FID))
#merge(P4,fid_pid,by.x="FID",by.y="FID") %>% select(PID) %>% table() %>% data.frame()
```

```
P5<-P3%>%filter(!is.na(FID))%>%merge(P4,fid_pid,by.x="FID",by.y="FID")
```

```

#mb %>% select(id) |> table() |> data.frame() |> filter(Freq==2) |> head(n=15)

#mb |> filter(id=="6006658")

#codes %>% filter(code2=="6006658")

#mb |> filter(id%in% c("6004319","6006658"))

mb2<-mb |> merge(fid_pid,by.x = "id",by.y="FID",all.x=TRUE)

mb3<-mb2 |> merge(codes,by.x="id",by.y="code1",all.x=TRUE) |>
  merge(codes,by.x="id",by.y="code2",all.x=TRUE) %>% mutate(acute=ifelse(!is.na(code2),"re
  mutate(rehab=ifelse(!is.na(code1),"acute","")) %>% mutate(phase=paste(acute,rehab,sep="")

#mb3|> filter(phase=="acute")

tfp<-fid_pid %>% select(PID) %>% table %>% data.frame() %>% arrange(-Freq)
#mb3 %>% filter(PID %in% tfp$PID)
#mb3 %>% filter(PID=="0152021")
#mb3 %>% select(PID) %>% table()%>%data.frame() %>% arrange(-Freq) %>% head(20)
#mb3 %>% select(PID,id,entry_date,phase)
ids<-mb3 |> filter(PID=="0019699") %>% select(id)
fids<-SCIM_17_20 %>% filter(FID %in% ids$id) %>% arrange(DATUM) %>% select(FID)

cpth<-readRDS("C:/Users/boris/Desktop/data hospital/cpth.rds")
ind_cpth <- read_excel("C:/Users/boris/Desktop/data hospital/ind_cpth.xlsx")
mb4<-mb3 %>% merge(cpth,by.x="id",by.y="V1") %>%
  merge(ind_cpth,by.x="clipath_code",by.y="Indication")

#setwd("C:/Users/boris/Desktop/SPZ hospital/SPZ-Project")

```

Dataset serv_cons

serv_cons: contains the aggregated information of FID and units of interventions:

```
#head(serv_cons)
```

```
time_cost<-readRDS("time_cost.rds")
serv_cons<-time_cost%>% select(-year) %>% group_by(id) %>%
  summarize(op_room=sum(op_room),anaest=sum(anaest),int_care=sum(int_care),imag=sum(imag),
```

Now, we identify the phases for every id in the previous table:

The first part merge by id and code1 (acute phase) to obtain uniq code and aggregate, the second part the same procedure but with code2 (rehab phase)

```
serv_cons1<-serv_cons%>% merge(codes%>% select(code1,uniq),by.x="id",by.y="code1") %>%
  merge(fid_pid,by.x="id",by.y="FID") %>% mutate(phase="acu")
```

```
serv_cons2<-serv_cons%>% merge(codes%>% select(code2,uniq),by.x="id",by.y="code2") %>%
  merge(fid_pid,by.x="id",by.y="FID") %>% mutate(phase="rehab")
```

```
serv_cons_phase<-rbind(serv_cons1,serv_cons2)
```

```
# We see the number of times that every FID appears in the table and phase
serv_cons_phase %>% select(PID) %>%
  table() %>% data.frame() %>% head() %>% arrange(-Freq) |> head()
```

	PID	Freq
1	0000374	6
2	0000362	4
3	0000477	2
4	0000511	2
5	0000548	2
6	0000567	2

```
# We check the first code
```

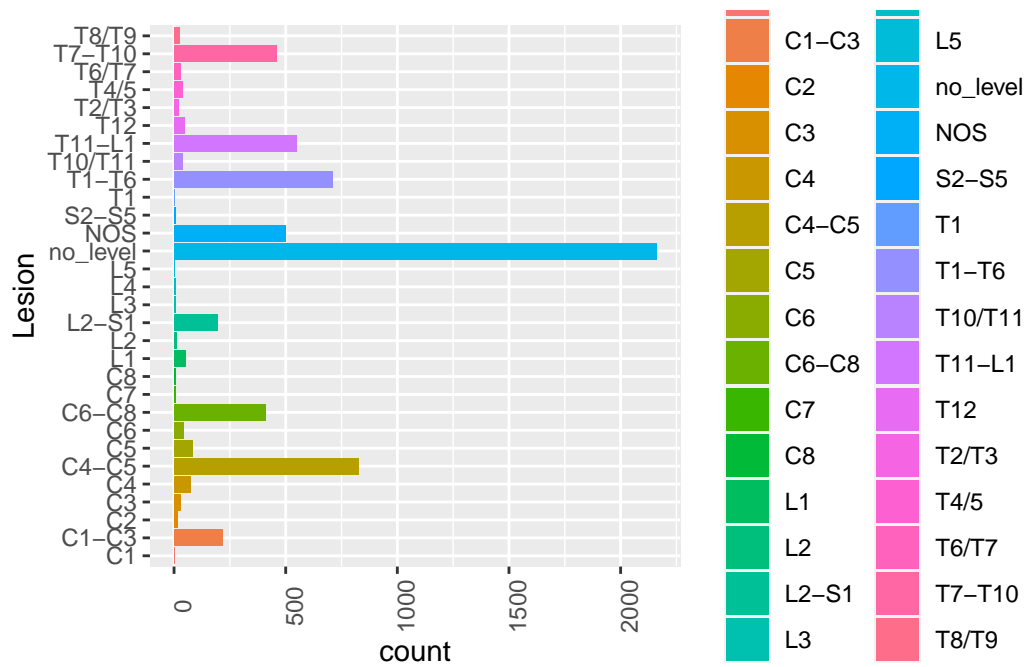
```
serv_cons_phase |> filter(PID=="0000374") %>% select(c(1,"PID",2,3,4,15,"uniq"))
```

	id	PID	op_room	anaest	int_care	phase	uniq
1	6028398	0000374	0	0	13398	acu	uni_3
2	6040865	0000374	152	478	287	acu	uni_4
3	6043203	0000374	0	0	555	acu	uni_5
4	6028919	0000374	0	0	0	rehab	uni_3

```
5 6041525 0000374      0      0      0 rehab uni_4
6 6043916 0000374      0      0     75 rehab uni_5
```

Lesion Level

```
mb %>% ggplot(aes(x=Lesion,fill=Lesion))+geom_bar(stat = "count")+theme(axis.text.x = element_text(angle=45))
```



```
mb_final<-mb |> merge(cpth,by.x="id",by.y="V1") |>merge(ind_cpth,by.x="clipath_code",by.y="V1")
```

```
mb_final$Lesion1<-substr(x = mb_final$Lesion,1,1)
```

```
#Threemap
```

```
tabtree<-mb_final |> select(main_diag,Lesion1)|> table()|> data.frame() |> group_by(main_d
```


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

```
#palet<-get_palette(palette="lancet",10)
#treemap(tabtree,index=c("Lesion1","main_diag"),vSize="tot",type="index",palette=palet)

mb_final$icddesc<-substr(mb_final$main_diag,1,3)

desc <-
  icd.data::icd10cm2016 |> filter(code%in% unique(mb_final$icddesc))|>select("code","short_desc")

mb_final<-mb_final|>merge(desc,by.x="icddesc",by.y="code")

tabtree<-mb_final |> select(Lesion1,short_desc)|> table()|> data.frame() |> group_by(Lesion1)
```

`summarise()` has grouped output by 'Lesion1'. You can override using the
`.groups` argument.

```
tabtree2<-tabtree|>filter(!short_desc %in% c("Paraplegia (paraparesis) and quadriplegia (quadriplegia)"))
#treemap(tabtree2,index=c("Lesion1","short_desc"),vSize="tot",type="index",palette=palet)

serv2<-serv_cons_phase|> merge(cpth,by.x="id",by.y="V1")|>merge(ind_cpth,by.x="clipath_code",by.y="V1")

Tab1<-mb_final %>% group_by(id)%>% summarize(length_stay=first(length_stay),gender=first(gender))

serv3<-serv2|>merge(Tab1,by.x="id",by.y="id")|>
  mutate(
    psy1=psyc/length_stay,
    op=op_room/length_stay,
    ana=anaest/length_stay,nurs=nurs/length_stay,int_care=int_care/length_stay,img=img/length_stay,
    summarize(
      avg_op=mean(op,na.rm=TRUE),avg_ana=mean(ana,na.rm=TRUE),avg_img=mean(img,na.rm=TRUE),
      avg_int=mean(int_care,na.rm=TRUE),avg_nurs=mean(nurs,na.rm=TRUE)
    )
  )

treemap_pathways<-gather(serv3,A,V,avg_op:avg_incare)

#treemap(treemap_pathways,index=c("Clinical.Pathway","A"),vSize="V",type="index",palette=palet)
```

```
treemap_pathways|>arrange(Clinical.Pathway,A,V)
```

```
# A tibble: 30 x 3
  Clinical.Pathway A           V
  <chr>           <chr>      <dbl>
1 Erstrehabilitation avg_ana    2.49
2 Erstrehabilitation avg_img    51.4
3 Erstrehabilitation avg_incare  4.94
4 Erstrehabilitation avg_lb     78.6
5 Erstrehabilitation avg_op     0.759
6 Handhandchirurgie avg_ana    37.3
7 Handhandchirurgie avg_img    23.2
8 Handhandchirurgie avg_incare  1.60
9 Handhandchirurgie avg_lb     49.0
10 Handhandchirurgie avg_op    12.0
# ... with 20 more rows
```

```
serv_ersrehab<-serv2|>merge(Tab1,by.x="id",by.y="id")|>
  mutate(psy1=psyc/length_stay,
         op=op_room/length_stay,
         ana=anaest/length_stay,nurs=nurs/length_stay,int_care=int_care/length_stay,img=img/length_stay)

serv_ersrehab|>group_by(gender)|> summarize(mean(psy1))
```

```
# A tibble: 2 x 2
  gender `mean(psy1)`
  <chr>      <dbl>
1 1          10.8
2 2          12.7
```

```
tab2<-serv2|>merge(Tab1,by.x="id",by.y="id")
tab22<-tab2 |> filter(Clinical.Pathway=="Erstrehabilitation",phase=="rehab") |> select(length_stay)

#plot(density(tab22$length_stay))
#ggbetweenstats(data = serv_ersrehab,x = gender,y=psy1)+scale_color_manual(values=colores)
```

```

#serv_ersrehab$Clinical.Pathway
#ggbetweenstats(data = serv_ersrehab,x = gender,y=op)+scale_color_manual(values=colores)

#ggbetweenstats(data = serv_ersrehab,x = gender,y=ana)+scale_color_manual(values=colores)

#ggbetweenstats(data = serv_ersrehab,x = gender,y=lb)+scale_color_manual(values=colores)

#ggbetweenstats(data = serv_ersrehab,x = gender,y=incare)+scale_color_manual(values=colores)

#ggbetweenstats(data = serv_ersrehab,x = gender,y=img)+scale_color_manual(values=colores)

#ggbetweenstats(data = serv_ersrehab,x = gender,y=nurs)+scale_color_manual(values=colores)


#ggplot(serv_ersrehab,aes(x=gender,y=psy1))+geom_boxplot()+  geom_jitter()+scale_color_manual(values=colores)

#apply(serv_ersrehab|>select(1:6),MARGIN = 2,FUN=function(x)mean(x,na.rm=TRUE))

```