

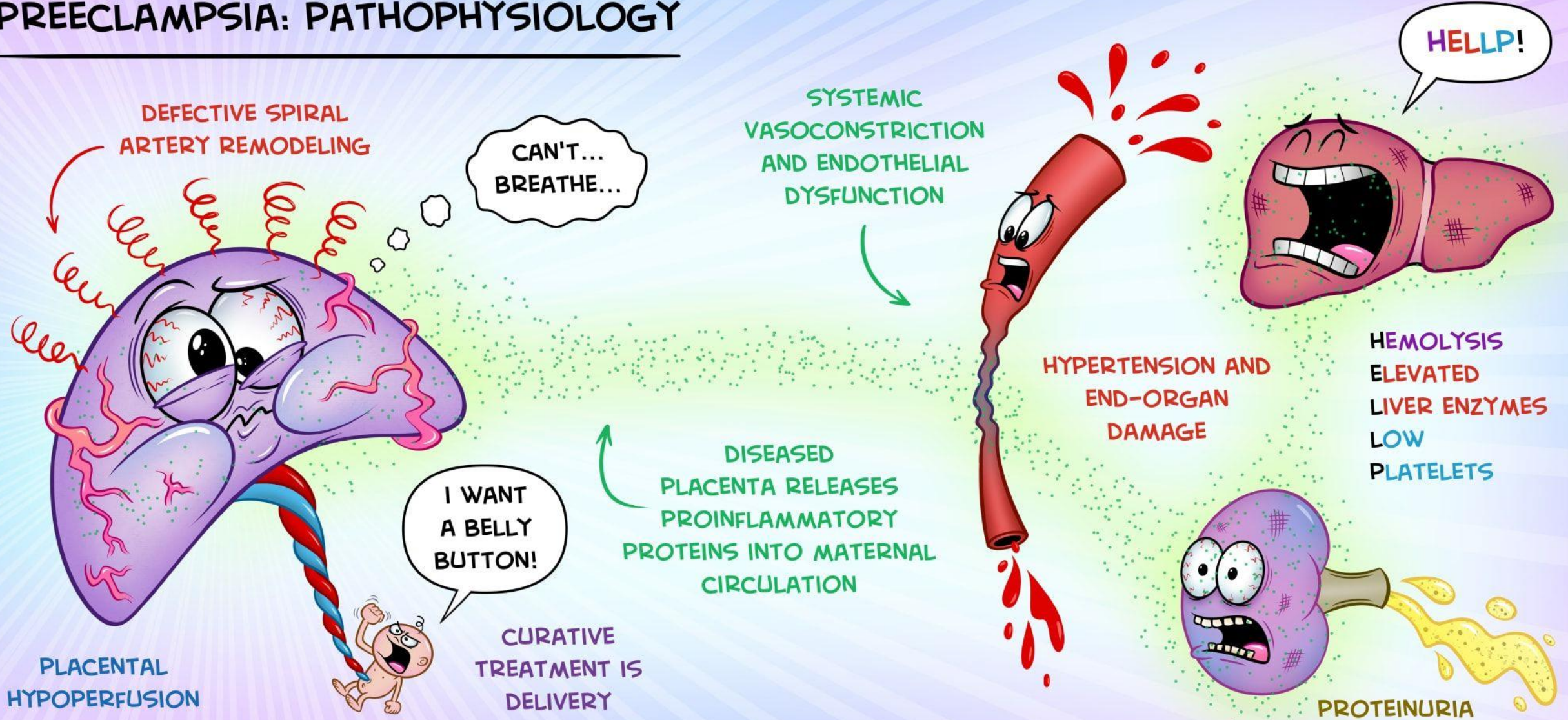
Control de calidad del cribado de preeclampsia

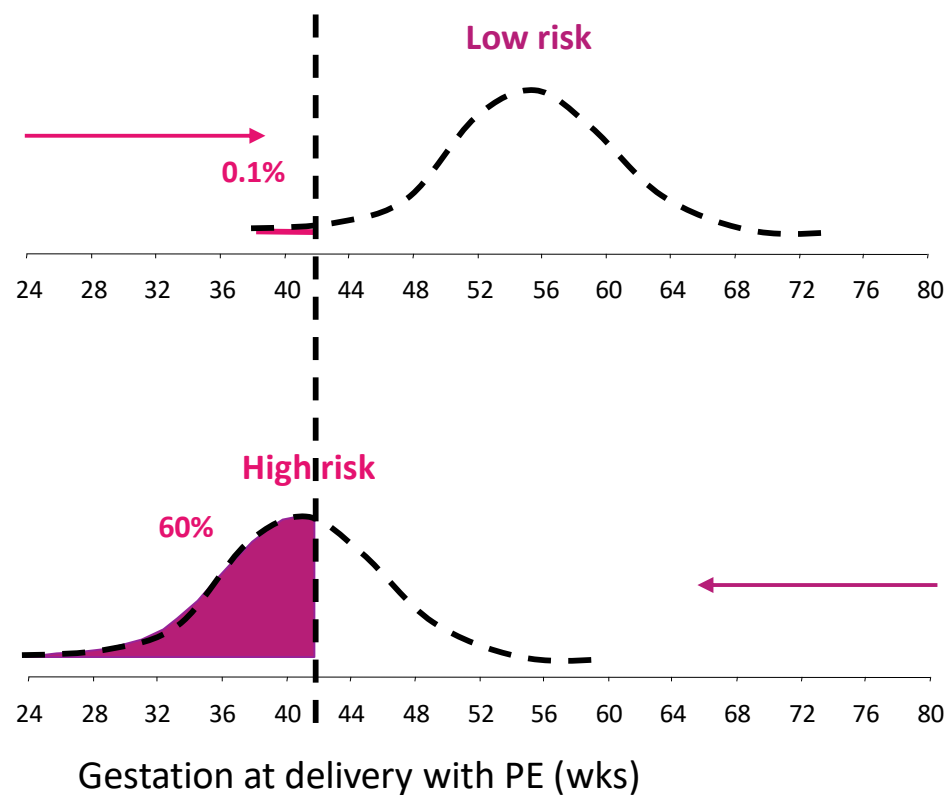
Análisis secundario del estudio PREVAL

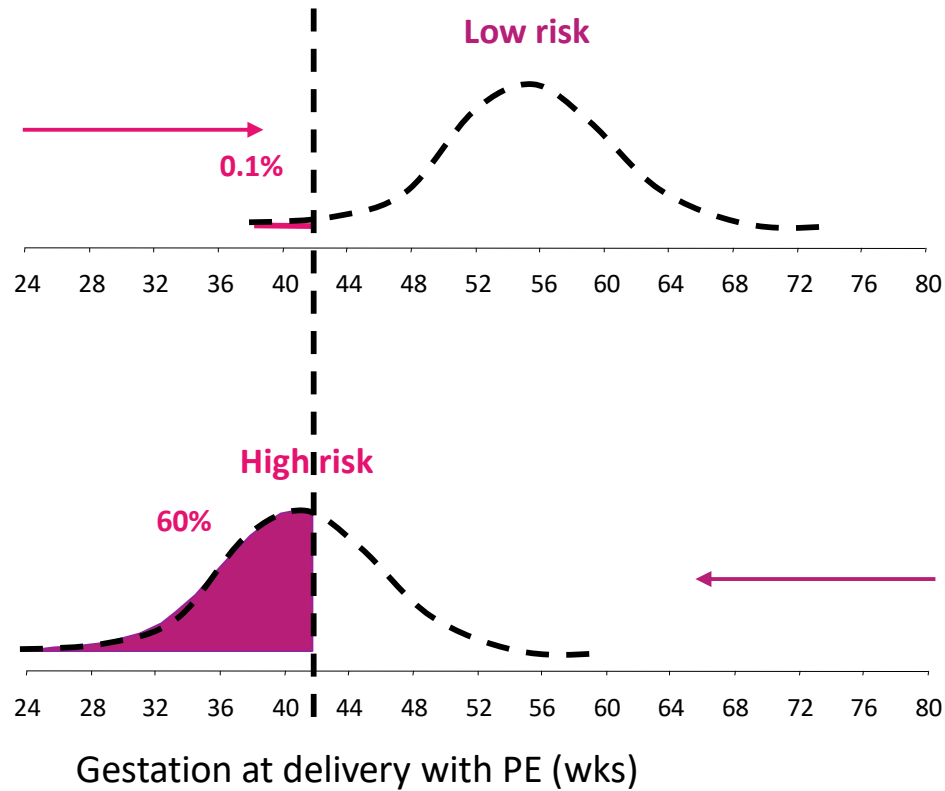
Valeria Rolle
Diana Cuenca
Katy de Paco
Nuria Valiño
Rocío Revello
Begoña Adiego
Belén Santacruz
María del Mar Gil



PREECLAMPSIA: PATHOPHYSIOLOGY







Age: every 10 years above 30 yrs
Weight: every 10 kg above 70 kg
Height: every 10 cm above 164 cm

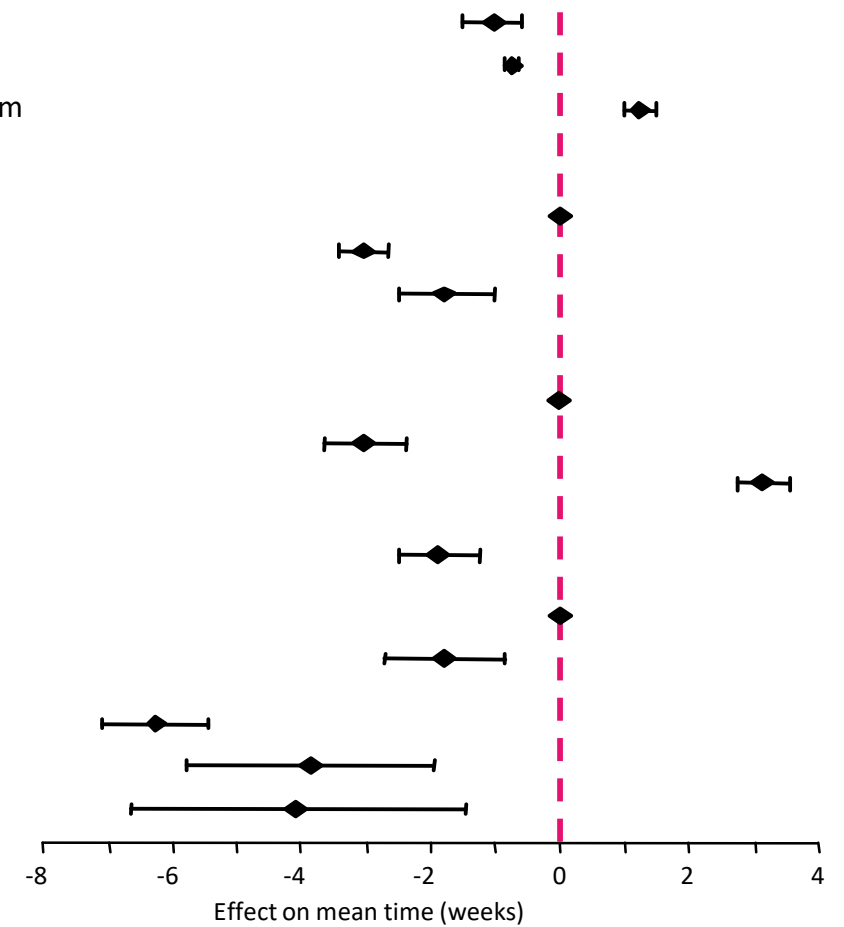
Racial origin
Caucasian
Afro-Caribbean
South Asian

Previous obstetric history
Nulliparous
Parous with preeclampsia
Parous with no preeclampsia

Mother had preeclampsia

Conception spontaneous
in vitro fertilization

Chronic hypertension
Type 1 diabetes mellitus
Systemic lupus erythematosus



Multicéntrico

Prospectivo

Se recogen:

- Factores maternos

- Presión arterial media

Poon 2012

**- Índice pulsatilidad arterias
uterinas**

Plasencia 2007

- PIGF

- PAPP-A

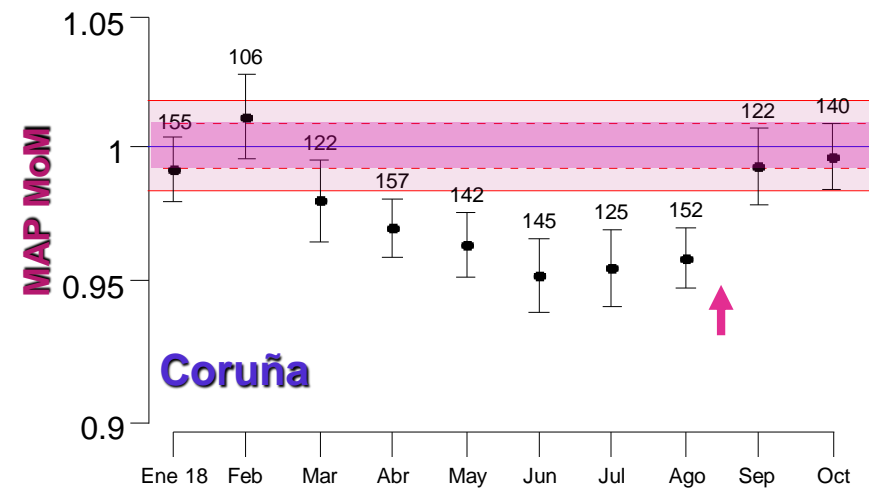
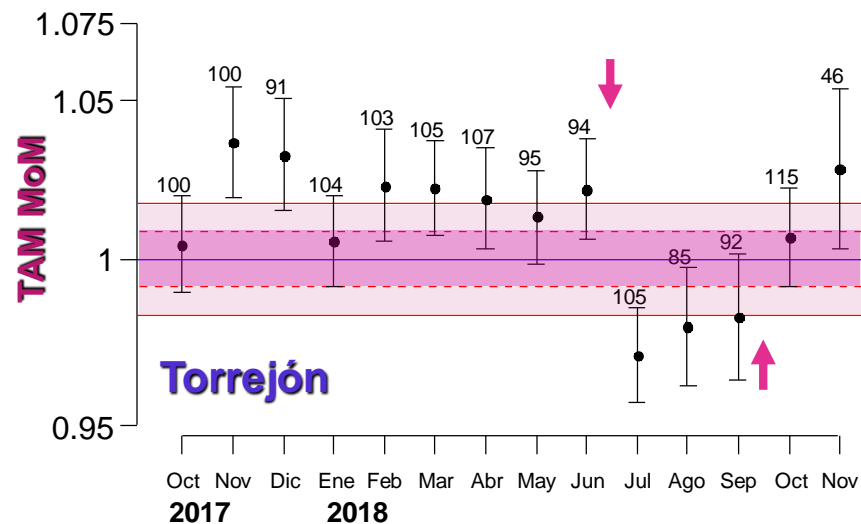


Características maternas; n = 5.874

Edad (años)	33,9 (30,1 – 36,9)
Peso (kg)	64,0 (57,2 – 73,1)
Altura (cm)	163 (159 – 167)
Edad gestacional (semanas)	12,7 (12,3 – 13,1)
Concepción espontánea	5,358 (91,2)
Nulíparas	2,986 (50,8)
Raza blanca	5,731 (97,6)
Fumadoras	716 (12,2)
LES / SAF	32 (0,5)
Diabetes mellitus	51 (0,9)
Hipertensión crónica	58 (1,0)
Toma de aspirina	295 (5%)

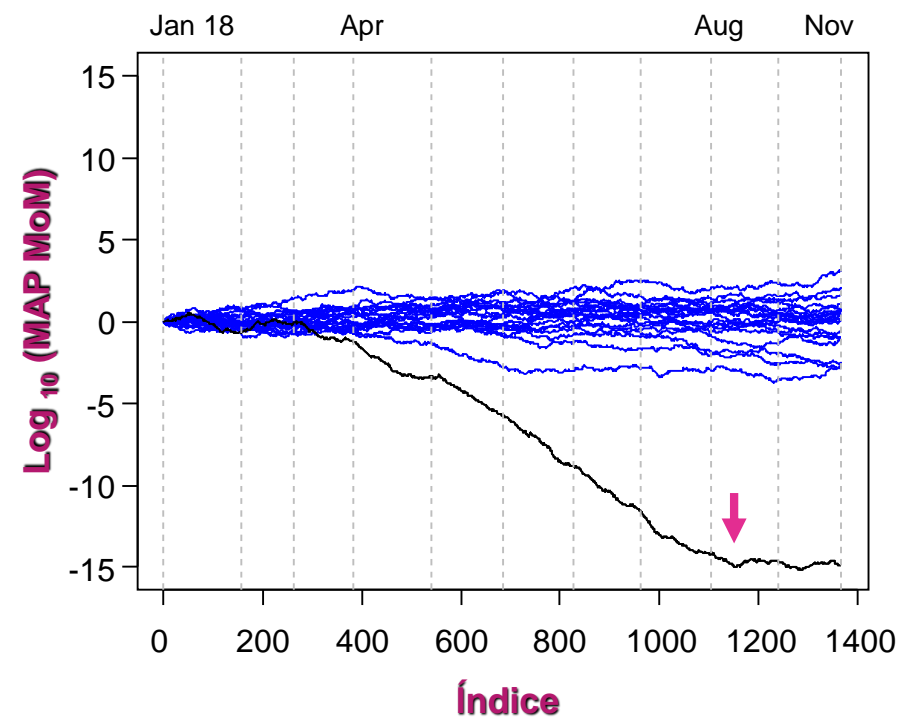
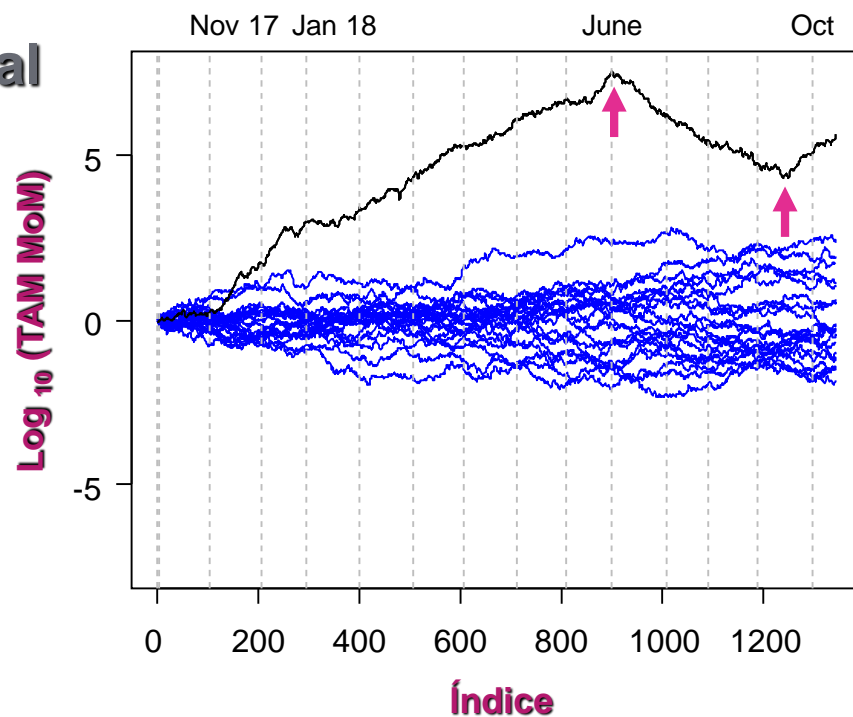
Mediana (RIQ) o N (%)



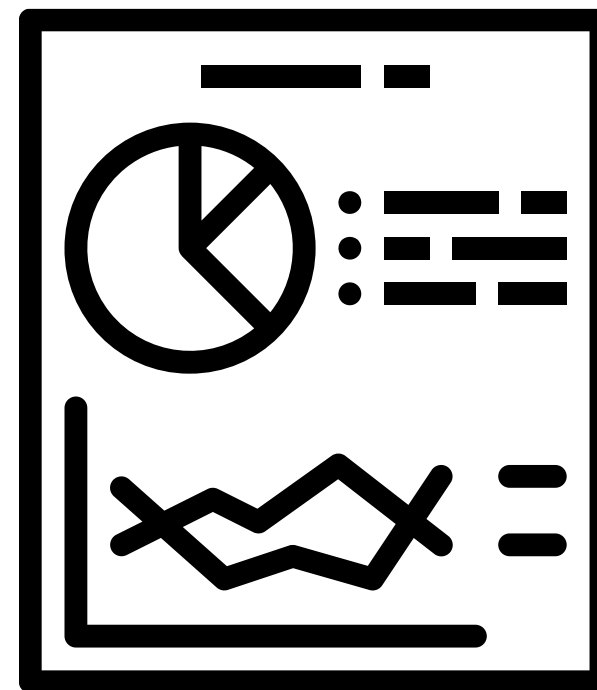
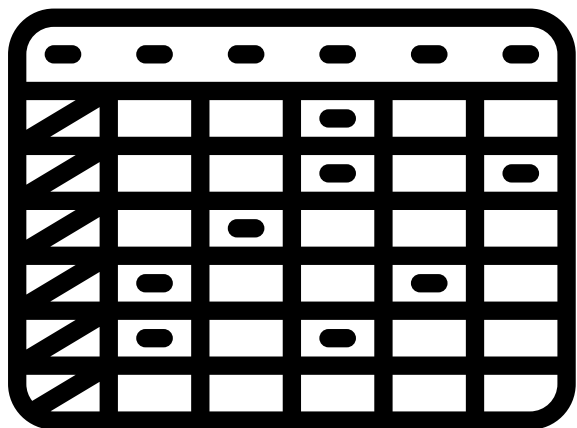


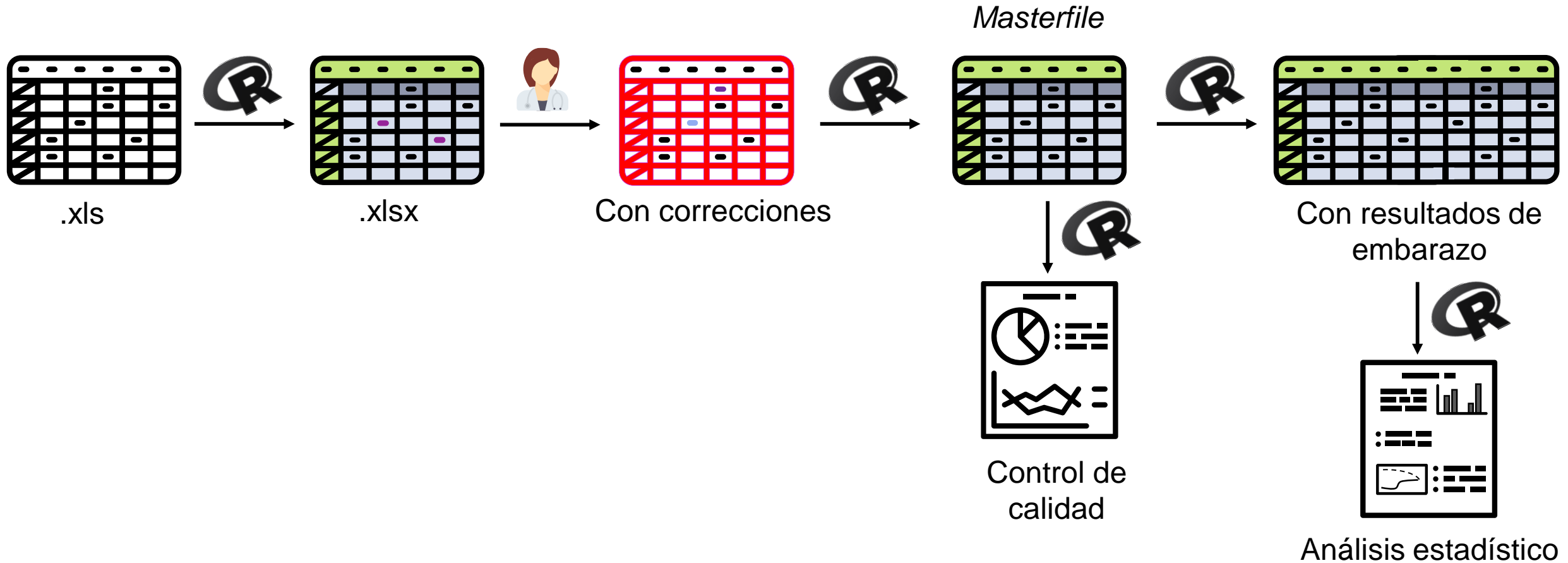
Necesidad de auditoría mensual:

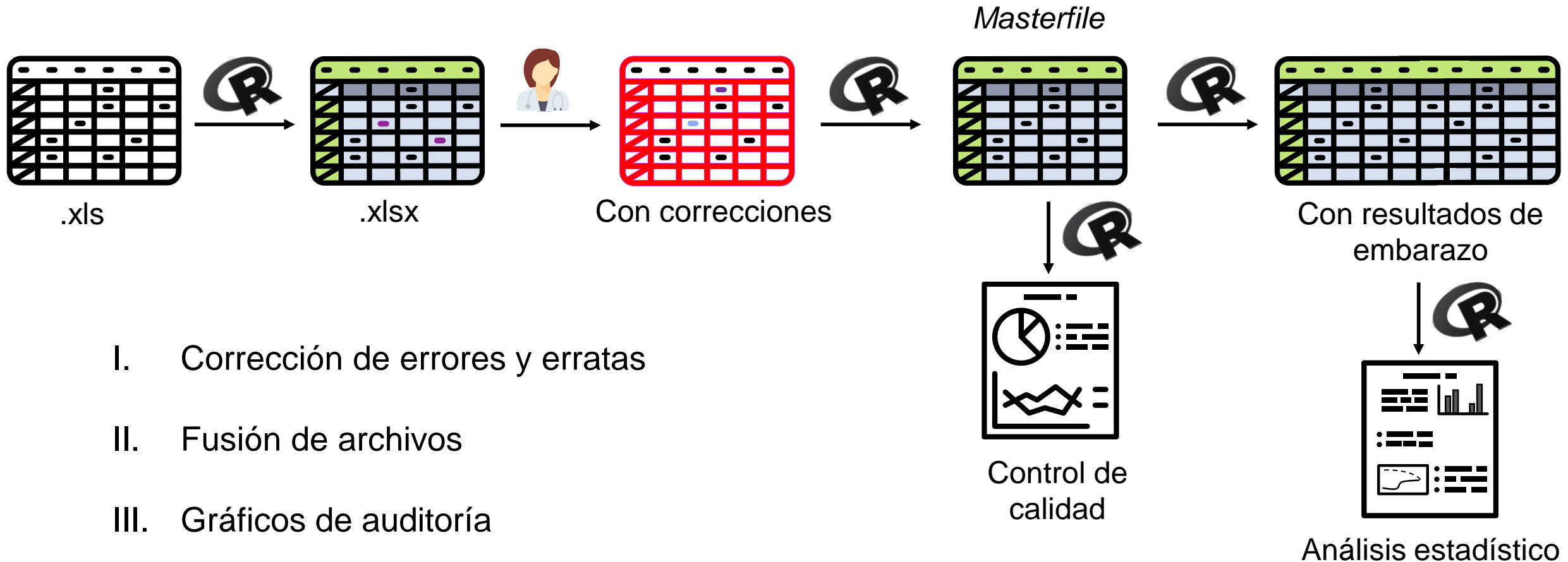
- Volver a entrenar personal
- Ajuste de MoMs

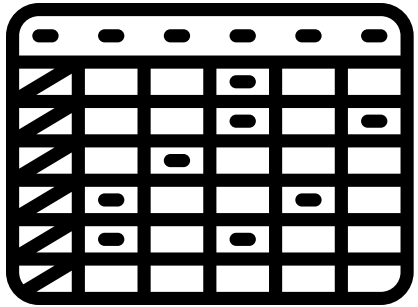


Situación ideal



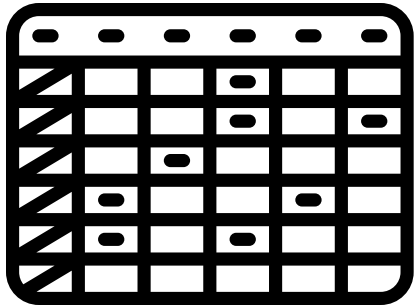






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```
dd1$com <- ifelse(dd1$peso > 150 | dd1$peso < 35, paste(dd1$com, "peso", sep = ", "), dd1$com)
dd1$com <- ifelse(dd1$altura > 215 | dd1$altura < 120, paste(dd1$com, "altura", sep = ", "), dd1$com)
dd1$com <- ifelse(is.na(dd1$dm), paste(dd1$com, "DM", sep = ", "), dd1$com)
dd1$com <- ifelse(is.na(dd1$hta), paste(dd1$com, "HTA", sep = ", "), dd1$com)
```

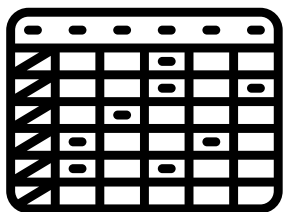


.xls

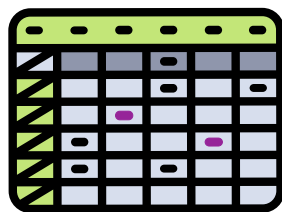
```
dd1$com <- ifelse((dd1$pe_ant_2 == "Nulip" &  
                  dd1$cir_previo == "Nulip" &  
                  dd1$paridad == "Nulip") &  
                  (dd1$emb1_pn != 171717.17 | dd1$emb2_pn != 171717.17),  
                  paste(dd1$com, "tiene info sobre emb anterior", sep = ", "),  
                  dd1$com)
```

Errores y erratas

Comentarios	Fecha_US	Case_N	DOB	Indicacion	Método	Concepcion	Ind_ovulac	Tabaco	Alcohol	Raza	Paridad	Drogas	DM
nulip con nacido vivo, ind. Ovula	3/5/18	2	1/2/92		ECO TA	espontáneo		no	no	Blanco	Nulip	no	no
	3/5/18	2	16/2/94	CPT	ECO TA	espontáneo	no	no	no	Negro	Nulip	no	no
	3/5/18	1	12/9/75	CPT	ECO TA	FIV	no	no	no	Blanco-Negro	Nulip	no	no
último parto en 1931	3/5/18	1	18/3/88	CPT	ECO TA	espontáneo	no	fumador	no	Blanco	Multip	no	no
raza, drogas	3/5/18	1	14/10/88	CPT	ECO TA	espontáneo	no	no	no		Nulip		no
	3/5/18	1	5/12/95	CPT	ECO TA	espontáneo	no	no	no	Negro	Nulip	no	no
	3/5/18	1	28/1/83	CPT	ECO TA	espontáneo	no	no	no	Blanco	Nulip	no	no
	3/5/18	1	18/9/87	CPT	ECO TA	espontáneo	no	no	no	Blanco	Nulip	no	no
	3/5/18	1	10/1/96	CPT	ECO TA	espontáneo	no	fumador	no	Blanco	Multip	no	no
ind. ovulación	3/5/18	1	10/5/78	CPT	ECO TA	FIV		no	no	Blanco	Multip	no	no
nulip con nacido vivo, ind. ovula	3/5/18	1	26/6/87	CPT	ECO TA	espontáneo		no	no	Blanco	Nulip	no	no
	3/5/18	1	29/1/88	CPT	ECO TA	espontáneo	no	fumador	no	Blanco	Nulip	no	no
tabaco, ind. ovulación	3/5/18	1	21/6/87	CPT		espontáneo		no - fumador	no	Blanco	Nulip	no	no
tabaco, ind. Ovulación	3/5/18	1	27/2/89	CPT		espontáneo		no - fumador	no	Blanco	Nulip	no	no
	4/5/18	1	24/6/85	CPT	ECO TA	espontáneo	no	no	no	Blanco	Multip	no	no
ind. ovulación	4/5/18	2	6/10/84			espontáneo		no	no	Blanco	Nulip	no	no
ind. ovulación	4/5/18	2	2/10/78		ECO TA	espontáneo		no	no	Blanco	Nulip	no	no
ind. ovulación	4/5/18	2	11/12/93			espontáneo		no	no	Blanco	Nulip	no	no
nulip con nacido vivo, ind. ovula	4/5/18	1	21/6/81			espontáneo		no	no	Blanco	Nulip	no	no
ind. ovulación	4/5/18	1	4/1/82		ECO TA Y T	espontáneo		no	no	Blanco	Nulip	no	no
	8/5/18	1	3/2/82	CPT	ECO TA	espontáneo	no	no	no	Blanco	Multip	no	no



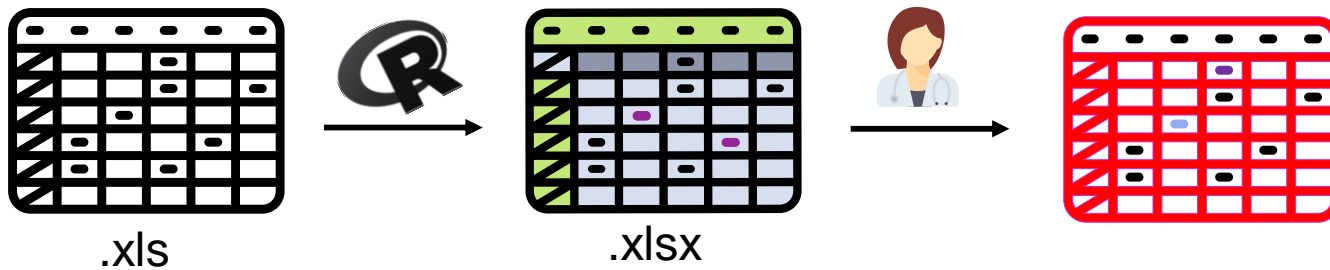
.xls



.xlsx



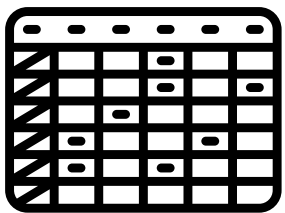
Raza	Freq.
Asian oriental	1
Asiático orienta	2
blanco	1
Blanco	2256
Blanco-Asiático oriental	3
Blanco-Negro	6
Asiático del sur	5
Mezcla	15
negro	1
Negro	32
Sudasiatico	1
Sudasiático	3



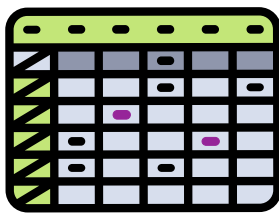
Raza	Freq.
Asian oriental	1
Asiático orienta	2
blanco	1
Blanco	2256
Blanco-Asiático oriental	3
Blanco-Negro	6
Asiático del sur	5
Mezcla	15
negro	1
Negro	32
Sudasiatico	1
Sudasiático	3

```
dd1$race <- car::recode(dd1$race, '
  "Mestizo"           = "Mixed";
  "Blanco-Negro"      = "Mixed";
  "Blanco-Asiatico oriental" = "Mixed";
  "Blanco-Sudasiático" = "Mixed";
  "Sudasiático-Asiático oriental" = "Mixed";
  "Negro-Asiático oriental" = "Mixed";
  "Asiático oriental"   = "East Asian";
  "Asiatico oriental"   = "East Asian";
  "Sudasiático"         = "South Asian";
  "Sudasiatico"         = "South Asian";
  "negro"               = "Black";
  "Negro"               = "Black";
  "blanco"              = "White";
  "Blanco"              = "White"
')
```

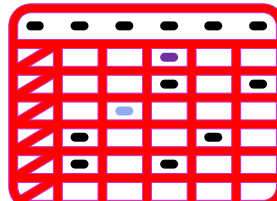
Raza	Freq.
Asian oriental	1
Black	33
East Asian	7
Mixed	6
South Asian	4
White	2257



.xls



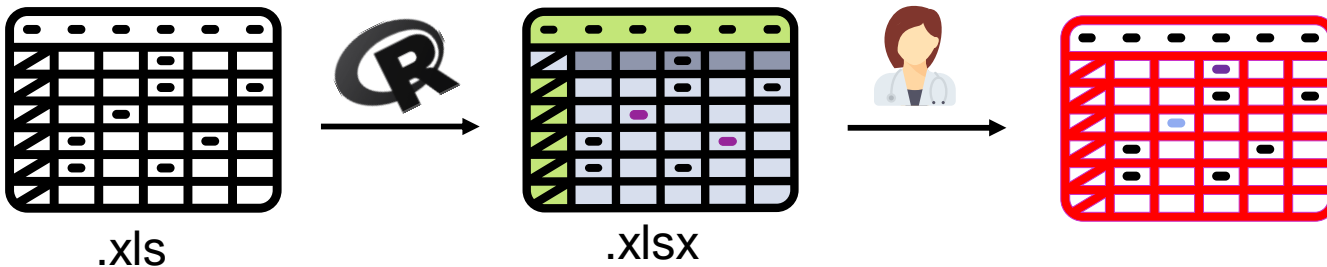
.xlsx



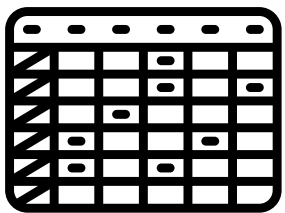
```
dd1$op <- tolower(dd$op)
```

```
dd1$op <- gsub("á", "a", dd1$op); dd1$op <- gsub("é", "e", dd1$op)  
dd1$op <- gsub("í", "i", dd1$op); dd1$op <- gsub("ó", "o", dd1$op)  
dd1$op <- gsub("ú", "u", dd1$op); dd1$op <- gsub(" ", "", dd1$op)
```

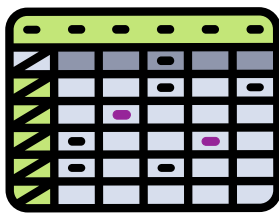
```
dd1$op <- car::recode(dd$op, '  
    "anamaria"      = "Ana María";  
    "luisacopon"    = "Luisa Copón")
```



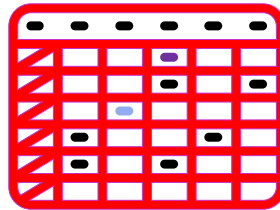
```
dd$m.operator <- gsub("MIR ", "", dd$m.operator)
dd$m.operator <- gsub("MIR - ", "", dd$m.operator)
dd$m.operator <- gsub(". MIR.", "", dd$m.operator)
dd$m.operator <- gsub(". MIR", "", dd$m.operator)
dd$m.operator <- gsub(" -MIR-", "", dd$m.operator)
dd$m.operator <- gsub(" _MIR_", "", dd$m.operator)
```



.xls

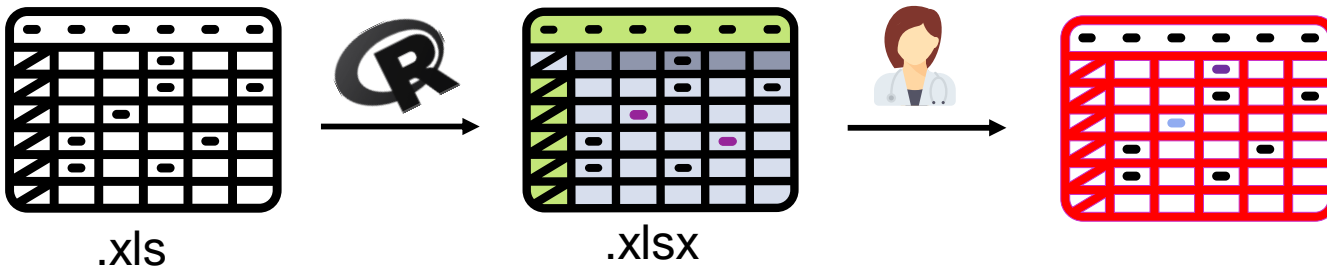


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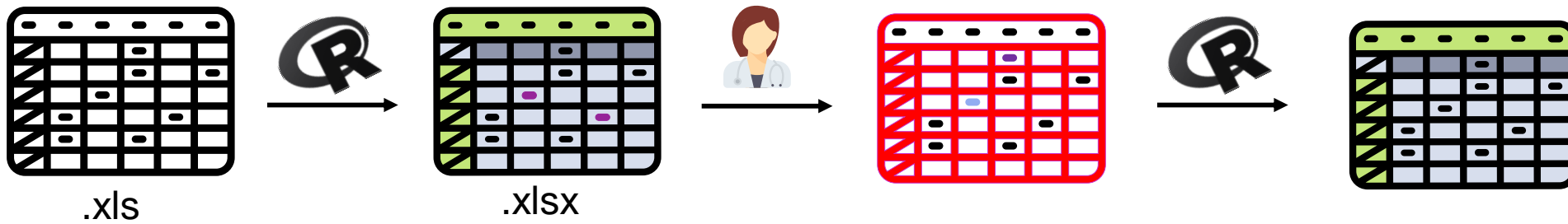


```
dd$m.operator <- gsub("MIR ", "", dd$m.operator)
dd$m.operator <- gsub("MIR - ", "", dd$m.operator)
dd$m.operator <- gsub(". MIR.", "", dd$m.operator)
dd$m.operator <- gsub(". MIR", "", dd$m.operator)
dd$m.operator <- gsub(" -MIR-", "", dd$m.operator)
dd$m.operator <- gsub(" _MIR_", "", dd$m.operator)
```

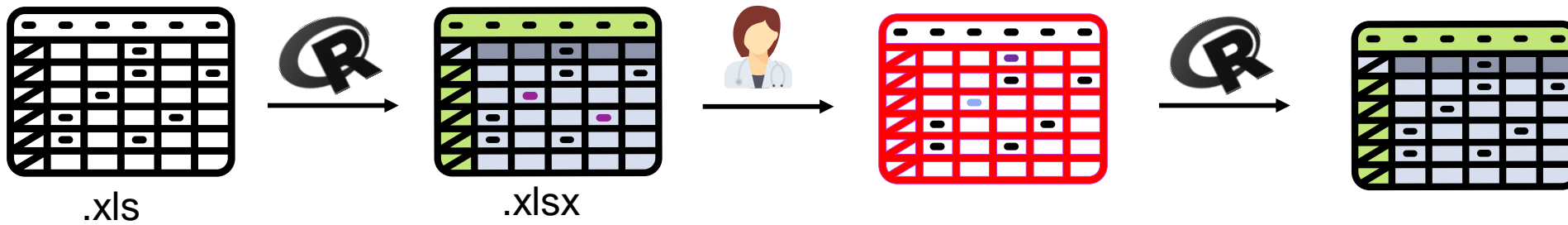
鶻	ú	桂	ñi
韃	ía	醬	án
鶻	ón	韓	ín
鶻	ós	韃	ía
鶻	á	醬	án
韃	íg	閏	ét
鶻	ár	鶻	óv



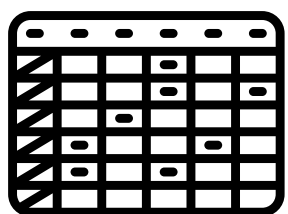
Fusión de archivos



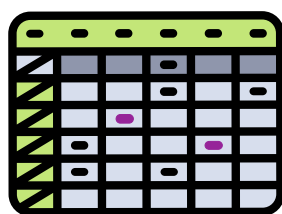
9 meses...



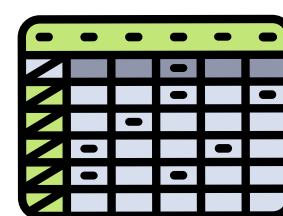
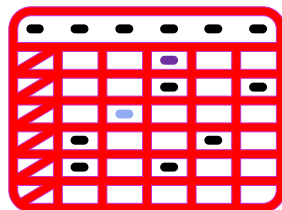
Fusión de archivos

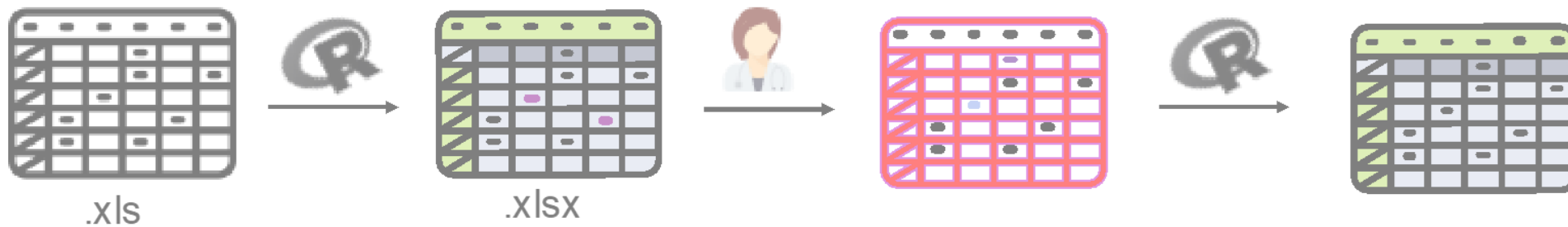
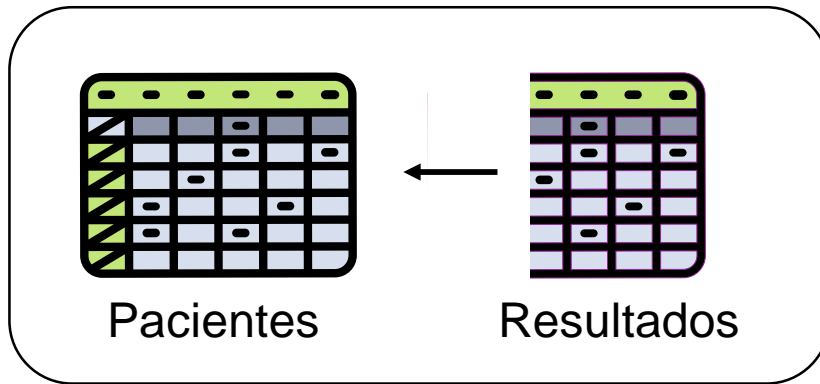


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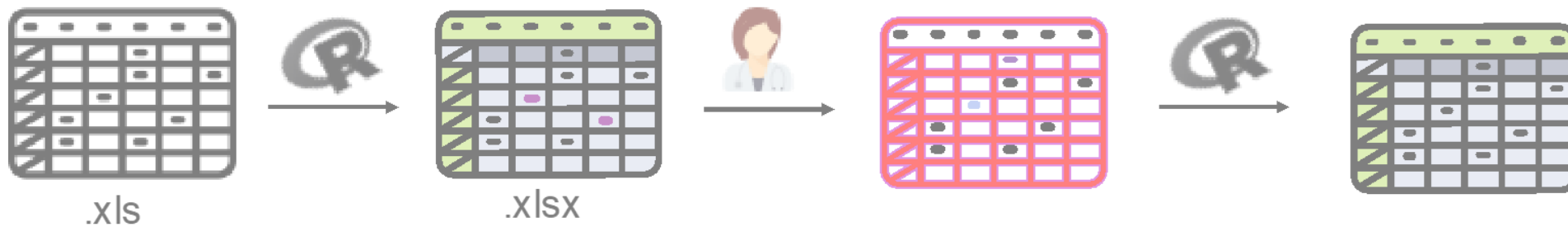
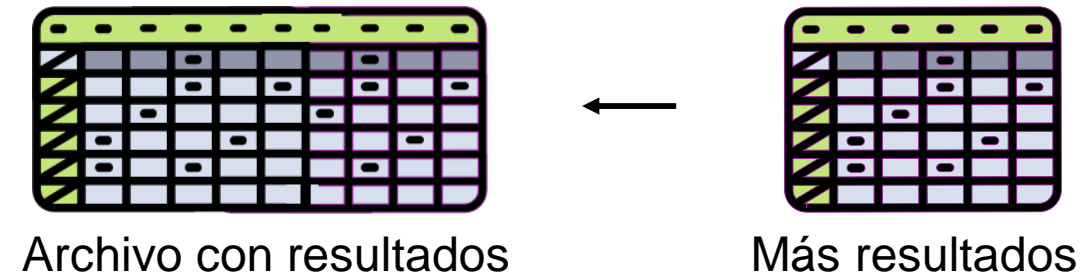
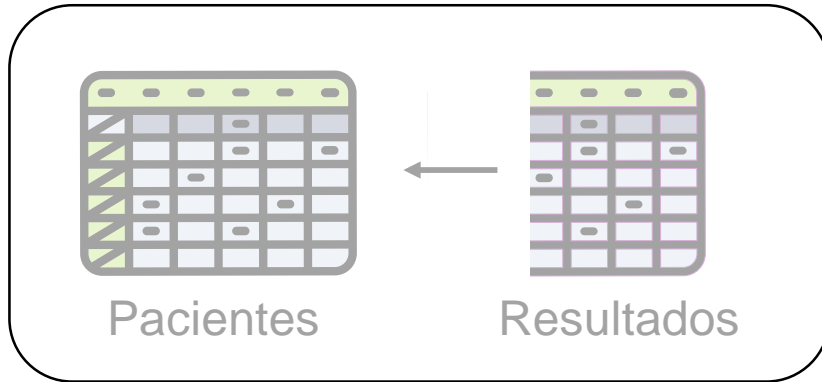


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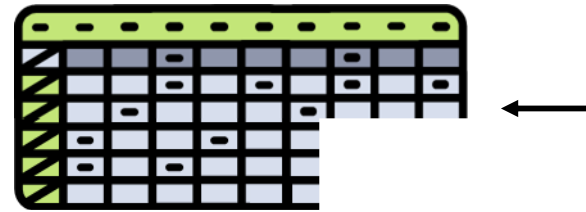




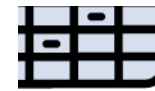
Fusión de archivos



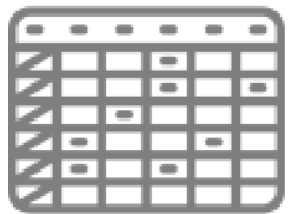
Fusión de archivos



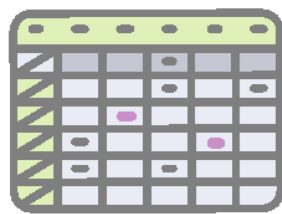
Archivo con resultados



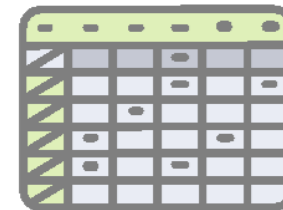
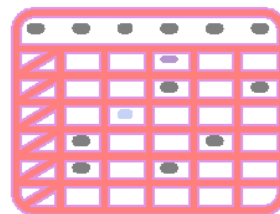
Más resultados y
actualizaciones



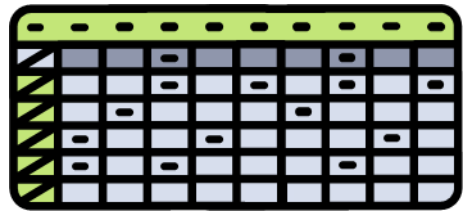
.xls



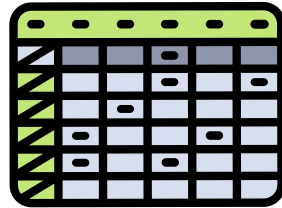
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Fusión de archivos

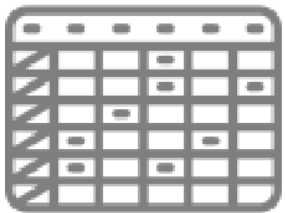


Archivo con resultados

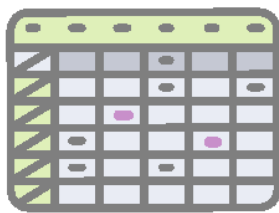


Más resultados y
actualizaciones

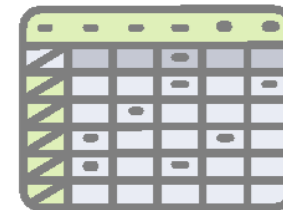
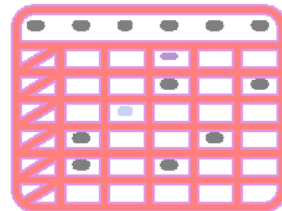
ID	NICU days.x	BW.x	Sex.x	NICU days.y	BW.y	Sex.y
14	0	2610	male	0	2610	male
40	0	342	female	0	3420	female
99	1	3040	female	1	3040	
102		1200	male	30	1200	male
103		3800	male	2	3800	male
107	0	2830	female	0	2830	female
141				0	3220	
155				0		male
173				0	3700	male

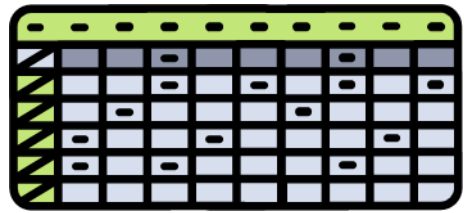


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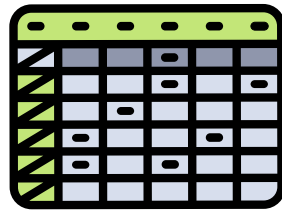


.xlsx





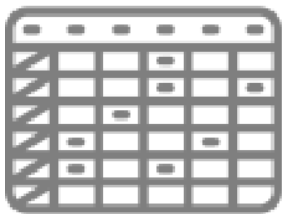
Archivo con resultados



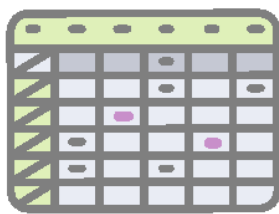
Más resultados y
actualizaciones

ID	NICU days.x	BW.x	Sex.x	NICU days.y	BW.y	Sex.y
14	0	2610	male	0	2610	male
40	0	342	female	0	3420	female
99	1	3040	female	1	3040	
102		1200	male	30	1200	male
103		3800	male	2	3800	male
107	0	2830	female	0	2830	female
141				0	3220	
155				0		male
173				0	3700	male

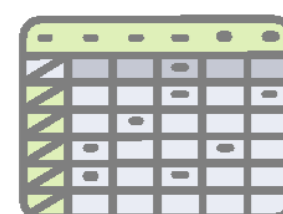
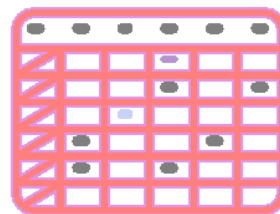
`dplyr::coalesce()`



.xls

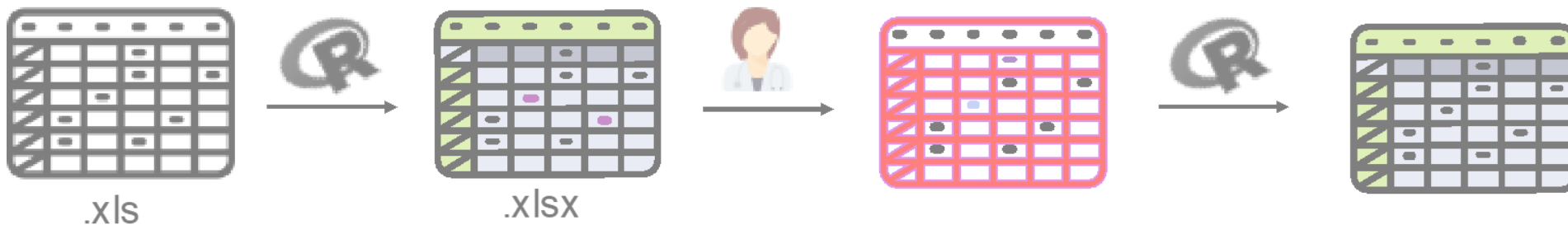


.xlsx



`dplyr::coalesce()`¹ + bucle o función

`rquery::natural_join()`²



¹ Hadley Wickham, Romain François, Lionel Henry and Kirill Müller (2019). *dplyr: A Grammar of Data Manipulation*. R package version 0.8.3. <https://CRAN.R-project.org/package=dplyr>

² John Mount (2019). *rquery: Relational Query Generator for Data Manipulation at Scale*. R package version 1.3.9. <https://CRAN.R-project.org/package=rquery>

`dplyr::coalesce()` ¹

```
coalesce_join <- function(x, y,
                          by = NULL, suffix = c(".x", ".y"),
                          join = dplyr::full_join, ...) {
  joined <- join(x, y, by = by, suffix = suffix, ...)
  # names of desired output
  cols <- union(names(x), names(y))

  to_coalesce <- names(joined)[!names(joined) %in% cols]
  suffix_used <- suffix[ifelse(endsWith(to_coalesce, suffix[1]), 1, 2)]
  # remove suffixes and deduplicate
  to_coalesce <- unique(substr(
    to_coalesce,
    1,
    nchar(to_coalesce) - nchar(suffix_used)
  ))

  coalesced <- purrr::map_dfc(to_coalesce, ~dplyr::coalesce(
    joined[[paste0(.x, suffix[1])]],
    joined[[paste0(.x, suffix[2])]]
  ))
  names(coalesced) <- to_coalesce

  dplyr::bind_cols(joined, coalesced)[cols]
}
```

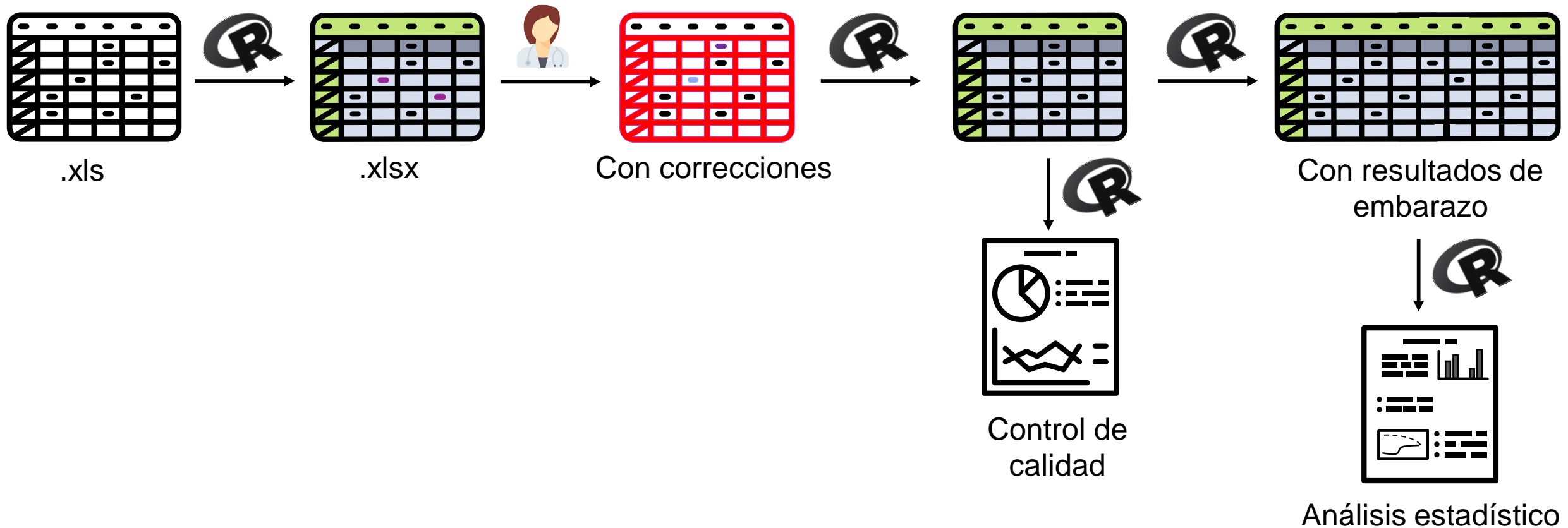
`dplyr::coalesce()` ¹

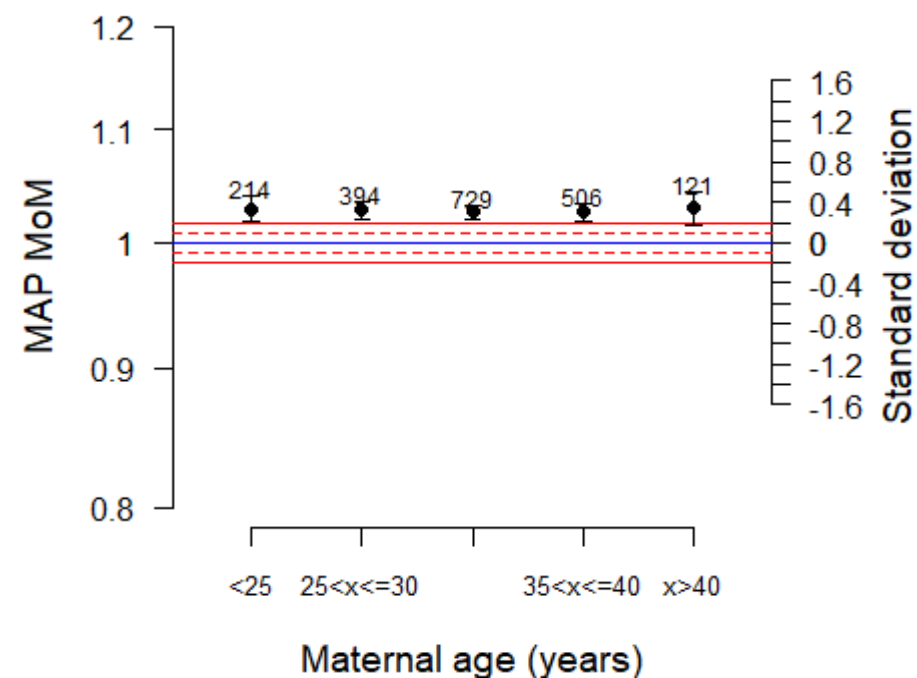
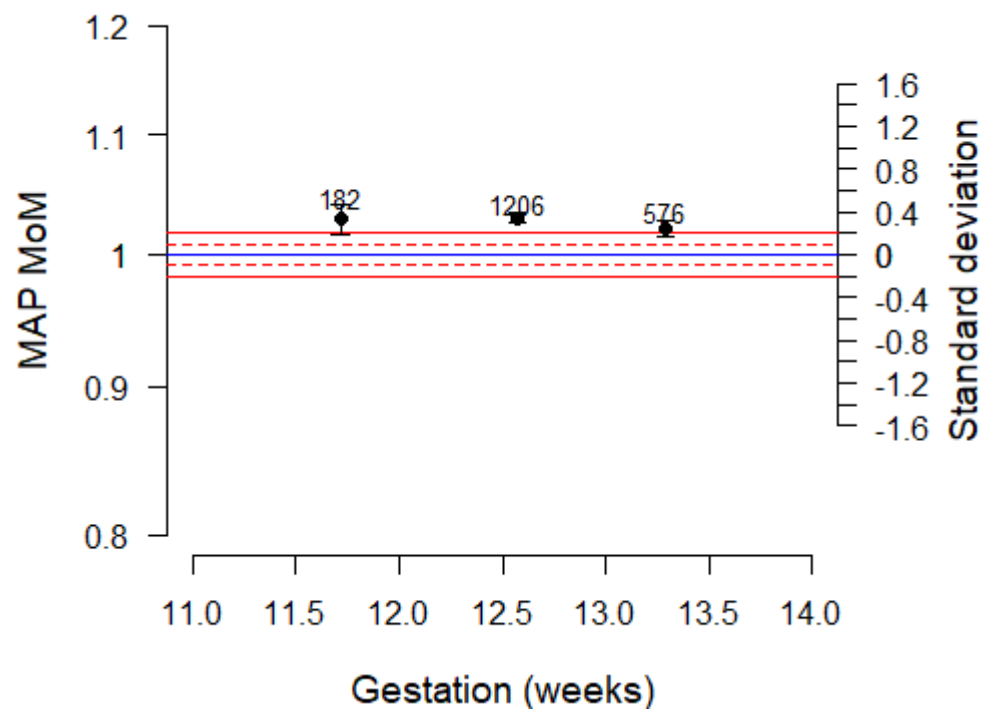
Merge + conseguir
nombres de columnas

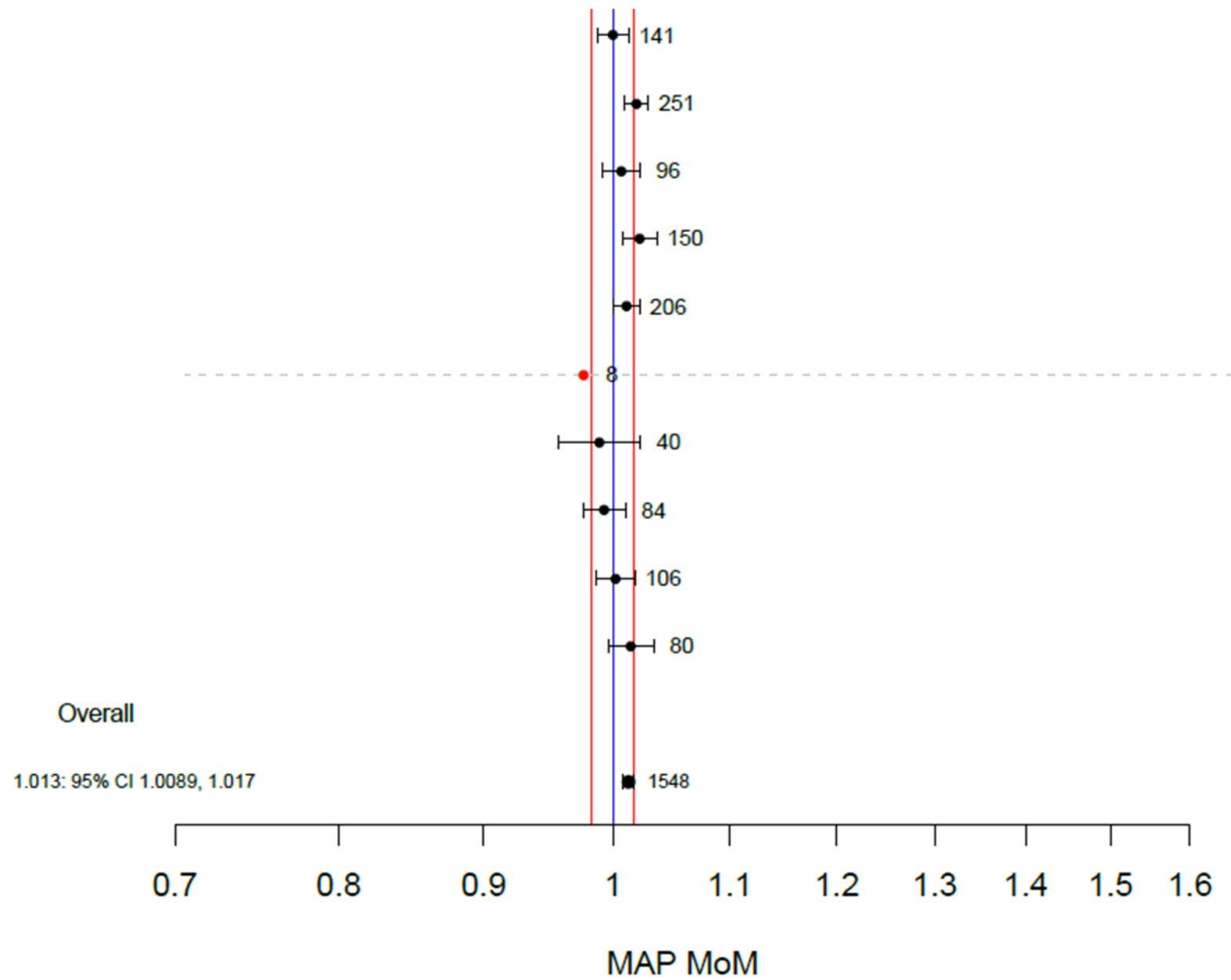
Coalesce de las
variables repetidas

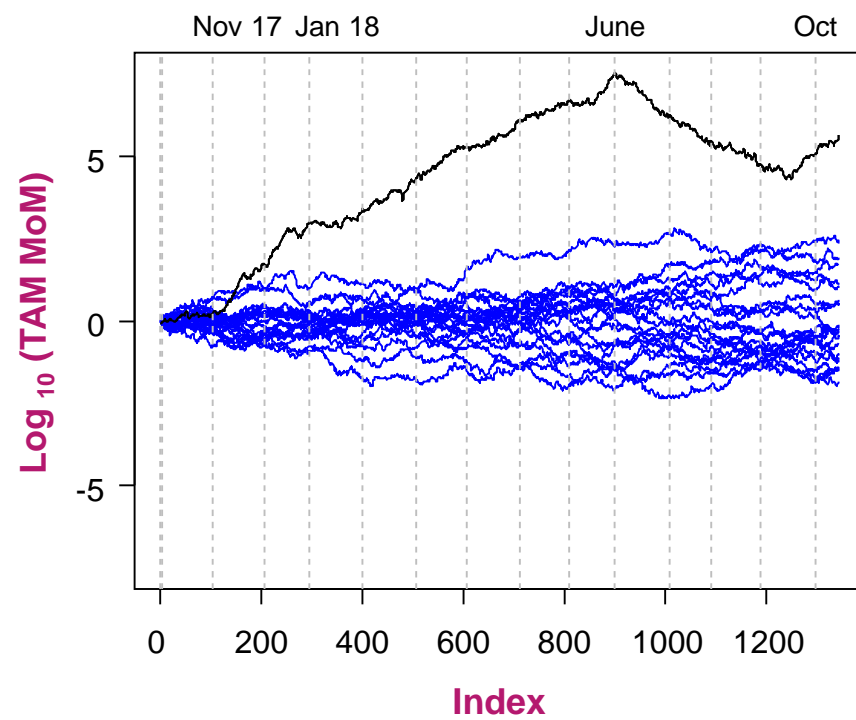
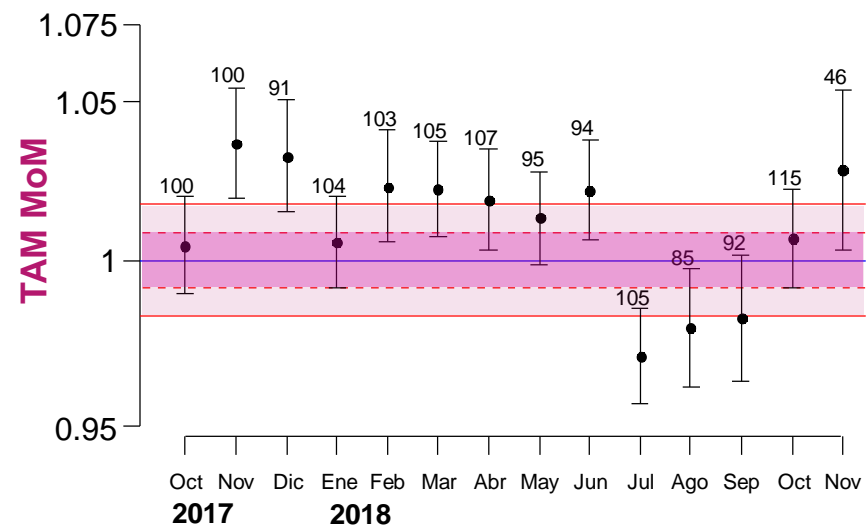
```
coalesce_join <- function(x, y,  
                          by = NULL, suffix = c(".x", ".y"),  
                          join = dplyr::full_join, ...) {  
  joined <- join(x, y, by = by, suffix = suffix, ...)  
  # names of desired output  
  cols <- union(names(x), names(y))  
  
  to_coalesce <- names(joined)[!names(joined) %in% cols]  
  suffix_used <- suffix[ifelse(endsWith(to_coalesce, suffix[1]), 1, 2)]  
  # remove suffixes and deduplicate  
  to_coalesce <- unique(substr(  
    to_coalesce,  
    1,  
    nchar(to_coalesce) - nchar(suffix_used)  
  ))  
  
  coalesced <- purrr::map_dfc(to_coalesce, ~dplyr::coalesce(  
    joined[[paste0(.x, suffix[1])]],  
    joined[[paste0(.x, suffix[2])]]  
  ))  
  names(coalesced) <- to_coalesce  
  
  dplyr::bind_cols(joined, coalesced)[cols]  
}
```


Fusión de archivos









- **Son necesarias auditorías periódicas y frecuentes.**
- **La auditoría de parámetros es efectiva en corregir desviaciones.**
- **R es una herramienta apropiada y eficaz para consolidar datos.**

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- Alan y David Wright por apoyo estadístico
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- Viewpoint y HNC por suministro del software

Gracias