

# Package: phoenix (via r-universe)

September 13, 2024

**Title** The Phoenix Pediatric Sepsis and Septic Shock Criteria

**Version** 1.1.1

**Description** Implementation of the Phoenix and Phoenix-8 Sepsis Criteria as described in ``Development and Validation of the Phoenix Criteria for Pediatric Sepsis and Septic Shock" by Sanchez-Pinto, Bennett, DeWitt, Russell et al. (2024) <[doi:10.1001/jama.2024.0196](https://doi.org/10.1001/jama.2024.0196)> (Drs. Sanchez-Pinto and Bennett contributed equally to this manuscript; Dr. DeWitt and Mr. Russell contributed equally to the manuscript), ``International Consensus Criteria for Pediatric Sepsis and Septic Shock" by Schlapbach, Watson, Sorce, Argent, et al. (2024) <[doi:10.1001/jama.2024.0179](https://doi.org/10.1001/jama.2024.0179)> (Drs Schlapbach, Watson, Sorce, and Argent contributed equally) and the application note ``phoenix: an R package and Python module for calculating the Phoenix pediatric sepsis score and criteria" by DeWitt, Russell, Rebull, Sanchez-Pinto, and Bennett (2024) <[doi:10.1093/jamiaopen/ooae066](https://doi.org/10.1093/jamiaopen/ooae066)>.

**Depends** R (>= 3.5.0)

**License** MIT + file LICENSE

**Encoding** UTF-8

**URL** <https://github.com/CU-DBMI-Peds/phoenix/>

**BugReports** <https://github.com/CU-DBMI-Peds/phoenix/issues>

**Language** en-us

**LazyData** true

**Suggests** knitr, ggplot2, qwraps2 (>= 0.6.0), reticulate, rmarkdown

**VignetteBuilder** knitr

**RoxygenNote** 7.3.2

**Repository** <https://cu-dbmi-peds.r-universe.dev>

**RemoteUrl** <https://github.com/cu-dbmi-peds/phoenix>

**RemoteRef** HEAD

**RemoteSha** 0c1d375cdc66ea91498b47aa410c3fbb41f6959a

## Contents

map . . . . .	2
phoenix . . . . .	3
phoenix8 . . . . .	6
phoenix_cardiovascular . . . . .	9
phoenix_coagulation . . . . .	12
phoenix_endocrine . . . . .	14
phoenix_hepatic . . . . .	15
phoenix_immunologic . . . . .	16
phoenix_neurologic . . . . .	18
phoenix_renal . . . . .	20
phoenix_respiratory . . . . .	22
sepsis . . . . .	24
<b>Index</b>	<b>26</b>

---

map	<i>Mean Arterial Pressure</i>
-----	-------------------------------

---

### Description

Estimate mean arterial pressure from systolic and diastolic blood pressures.

### Usage

```
map(sbp, dbp)
```

### Arguments

sbp	numeric vector, systolic blood pressure measured in mmHg
dbp	numeric vector, diastolic blood pressure measured in mmHg

### Details

Mean Arterial Pressure is approximated by:  $(DBP + (SBP - DBP) / 3) = (2/3) DBP + (1/3) SBP$

### Value

a numeric vector

**Examples**

```

DF <- expand.grid(sbp = 40:130, dbp = 20:100)
DF$map <- with(DF, map(sbp, dbp))
with(DF, plot(sbp, dbp, col = map))
DF$map[DF$sbp < DF$dbp] <- NA

z <- matrix(DF$map, nrow = length(unique(DF$sbp)), ncol = length(unique(DF$dbp)))

image(
  x = unique(DF$sbp),
  y = unique(DF$dbp),
  z = z,
  col = hcl.colors(100, palette = "RdBu"),
  xlab = "SBP (mmHg)",
  ylab = "DBP (mmHg)",
  main = "Estimated Mean Arterial Pressue"
)
contour(x = unique(DF$sbp), y = unique(DF$dbp), z = z, add = TRUE)

```

---

 phoenix

*The Phoenix Sepsis Score*


---

**Description**

The diagnostic Phoenix Sepsis Criteria based on four organ dysfunction scores, respiratory, cardiovascular, coagulation, and neurologic. A score of 2 or more indicates sepsis.

**Usage**

```

phoenix(
  pf_ratio,
  sf_ratio,
  imv,
  other_respiratory_support,
  vasoactives,
  lactate,
  map,
  platelets,
  inr,
  d_dimer,
  fibrinogen,
  gcs,
  fixed_pupils,
  age,
  data = parent.frame(),
  ...
)

```

**Arguments**

<code>pf_ratio</code>	numeric vector for the PaO <sub>2</sub> /FiO <sub>2</sub> ratio; PaO <sub>2</sub> = arterial oxygen pressure; FiO <sub>2</sub> = fraction of inspired oxygen; PaO <sub>2</sub> is measured in mmHg and FiO <sub>2</sub> is from 0.21 (room air) to 1.00.
<code>sf_ratio</code>	numeric vector for the SpO <sub>2</sub> /FiO <sub>2</sub> ratio; SpO <sub>2</sub> = oxygen saturation, measured in a percent; ratio for 92% oxygen saturation on room air is 92/0.21 = 438.0952.
<code>imv</code>	invasive mechanical ventilation; numeric or integer vector, (0 = not intubated; 1 = intubated)
<code>other_respiratory_support</code>	other respiratory support; numeric or integer vector, (0 = no support; 1 = support)
<code>vasoactives</code>	an integer vector, the number of systemic vasoactive medications being administered to the patient. Six vasoactive medications are considered: dobutamine, dopamine, epinephrine, milrinone, norepinephrine, vasopressin.
<code>lactate</code>	numeric vector with the lactate value in mmol/L
<code>map</code>	numeric vector, mean arterial pressure in mmHg
<code>platelets</code>	numeric vector for platelets counts in units of 1,000/uL (thousand per microliter)
<code>inr</code>	numeric vector for the international normalised ratio blood test
<code>d_dimer</code>	numeric vector for D-Dimer, units of mg/L FEU
<code>fibrinogen</code>	numeric vector units of mg/dL
<code>gcs</code>	integer vector; total Glasgow Comma Score
<code>fixed_pupils</code>	integer vector; 1 = bilaterally fixed pupil, 0 = otherwise
<code>age</code>	numeric vector age in months
<code>data</code>	a list, data.frame, or environment containing the input vectors
<code>...</code>	pass through

**Details**

The details of each of the four component scores are found in there respective help files.

**Value**

A data.frame with seven columns:

1. `phoenix_respiratory_score`
2. `phoenix_cardiovascular_score`
3. `phoenix_coagulation_score`
4. `phoenix_neurologic_score`
5. `phoenix_sepsis_score`
6. `phoenix_sepsis` An integer vector, 0 = not septic, 1 = septic (score greater or equal to 2)
7. `phoenix_septic_shock` An integer vector, 0 = not septic shock, 1 = septic shock (score greater or equal 2 and cardiovascular dysfunction)

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

## References

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

## See Also

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

## Examples

```
# Using the example sepsis data set, read more details in the vignette
phoenix_scores <-
  phoenix(
    # respiratory
    pf_ratio = pao2 / fio2,
    sf_ratio = ifelse(spo2 <= 97, spo2 / fio2, NA_real_),
    imv = vent,
    other_respiratory_support = as.integer(fio2 > 0.21),
    # cardiovascular
    vasoactives = dobutamine + dopamine + epinephrine + milrinone + norepinephrine + vasopressin,
    lactate = lactate,
    age = age,
    map = dbp + (sbp - dbp)/3,
    # coagulation
    platelets = platelets,
    inr = inr,
    d_dimer = d_dimer,
    fibrinogen = fibrinogen,
    # neurologic
    gcs = gcs_total,
    fixed_pupils = as.integer(pupil == "both-fixed"),
    data = sepsis
  )

str(phoenix_scores)
```

---

 phoenix8

*The Phoenix 8 Sepsis Score*


---

### Description

The extended Phoenix criteria using a total eight organ systems. This is intended mostly for research as an extension of the Phoenix Sepsis Criteria which is based on four organ systems.

### Usage

```

phoenix8(
  pf_ratio,
  sf_ratio,
  imv,
  other_respiratory_support,
  vasoactives,
  lactate,
  map,
  platelets,
  inr,
  d_dimer,
  fibrinogen,
  gcs,
  fixed_pupils,
  glucose,
  anc,
  alc,
  creatinine,
  bilirubin,
  alt,
  age,
  data = parent.frame(),
  ...
)

```

### Arguments

pf_ratio	numeric vector for the PaO <sub>2</sub> /FiO <sub>2</sub> ratio; PaO <sub>2</sub> = arterial oxygen pressure; FiO <sub>2</sub> = fraction of inspired oxygen; PaO <sub>2</sub> is measured in mmHg and FiO <sub>2</sub> is from 0.21 (room air) to 1.00.
sf_ratio	numeric vector for the SpO <sub>2</sub> /FiO <sub>2</sub> ratio; SpO <sub>2</sub> = oxygen saturation, measured in a percent; ratio for 92% oxygen saturation on room air is 92/0.21 = 438.0952.
imv	invasive mechanical ventilation; numeric or integer vector, (0 = not intubated; 1 = intubated)
other_respiratory_support	other respiratory support; numeric or integer vector, (0 = no support; 1 = support)

vasoactives	an integer vector, the number of systemic vasoactive medications being administered to the patient. Six vasoactive medications are considered: dobutamine, dopamine, epinephrine, milrinone, norepinephrine, vasopressin.
lactate	numeric vector with the lactate value in mmol/L
map	numeric vector, mean arterial pressure in mmHg
platelets	numeric vector for platelets counts in units of 1,000/uL (thousand per microliter)
inr	numeric vector for the international normalised ratio blood test
d_dimer	numeric vector for D-Dimer, units of mg/L FEU
fibrinogen	numeric vector units of mg/dL
gcs	integer vector; total Glasgow Comma Score
fixed_pupils	integer vector; 1 = bilaterally fixed pupil, 0 = otherwise
glucose	numeric vector; blood glucose measured in mg/dL
anc	absolute neutrophil count; a numeric vector; units of 1,000 cells per cubic millimeter
alc	absolute lymphocyte count; a numeric vector; units of 1,000 cells per cubic millimeter
creatinine	numeric vector; units of mg/dL
bilirubin	numeric vector; units of mg/dL
alt	alanine aminotransferase; a numeric vector; units of IU/L
age	numeric vector age in months
data	a list, data.frame, or environment containing the input vectors
...	pass through

### Details

The Phoenix Sepsis Criteria is based on the score form respiratory, cardiovascular, coagulation, and neurologic. Phoenix 8 uses these four an endocrine, immunologic, renal, and hepatic. Details on the scoring for each of the eight component organ systems are found in the respective manual files.

### Value

a data.frame with 12 integer columns.

1. phoenix\_respiratory\_score
2. phoenix\_cardiovascular\_score
3. phoenix\_coagulation\_score
4. phoenix\_neurologic\_score
5. phoenix\_sepsis\_score
6. phoenix\_sepsis 0 = not septic; 1 = septic (phoenix\_sepsis\_score greater or equal 2)
7. phoenix\_septic\_shock 0 = no septic shock; 1 = septic shock (sepsis with cardiovascular dysfunction)
8. phoenix\_endocrine\_score

9. phoenix\_immunologic\_score
10. phoenix\_renal\_score
11. phoenix\_hepatic\_score
12. phoenix8\_sepsis\_score

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

## References

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

## See Also

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

## Examples

```
# Using the example sepsis data set, read more details in the vignette
phoenix8_scores <-
  phoenix8(
    # respiratory
    pf_ratio = pao2 / fio2,
    sf_ratio = ifelse(spo2 <= 97, spo2 / fio2, NA_real_),
    imv = vent,
    other_respiratory_support = as.integer(fio2 > 0.21),
    # cardiovascular
    vasoactives = dobutamine + dopamine + epinephrine + milrinone + norepinephrine + vasopressin,
    lactate = lactate,
    age = age, # Also used in the renal assessment.
    map = dbp + (sbp - dbp)/3,
    # coagulation
    platelets = platelets,
    inr = inr,
    d_dimer = d_dimer,
    fibrinogen = fibrinogen,
    # neurologic
```



```

      gcs = gcs_total,
      fixed_pupils = as.integer(pupil == "both-fixed"),
# endocrine
      glucose = glucose,
# immunologic
      anc = anc,
      alc = alc,
# renal
      creatinine = creatinine,
# no need to specify age again
# hepatic
      bilirubin = bilirubin,
      alt = alt,
      data = sepsis
    )

str(phoenix8_scores)

```

---

 phoenix\_cardiovascular

*Phoenix Cardiovascular Score*

---

## Description

Generate the cardiovascular organ system dysfunction score as part of the diagnostic Phoenix Sepsis Criteria.

## Usage

```

phoenix_cardiovascular(
  vasoactives = NA_integer_,
  lactate = NA_real_,
  age = NA_real_,
  map = NA_real_,
  data = parent.frame(),
  ...
)

```

## Arguments

vasoactives	an integer vector, the number of systemic vasoactive medications being administered to the patient. Six vasoactive medications are considered: dobutamine, dopamine, epinephrine, milrinone, norepinephrine, vasopressin.
lactate	numeric vector with the lactate value in mmol/L
age	numeric vector age in months
map	numeric vector, mean arterial pressure in mmHg
data	a list, data.frame, or environment containing the input vectors
...	pass through

**Details**

There were six systemic vasoactive medications considered when the Phoenix criteria was developed: dobutamine, dopamine, epinephrine, milrinone, norepinephrine, and vasopressin.

During development, the values used for map were taken preferentially from arterial measurement, then cuff measures, and provided values before approximating the map from blood pressure values via  $DBP + 1/3 (SBP - DBP)$ , where DBP is the diastolic blood pressure and SBP is the systolic blood pressure.

**Value**

a integer vector with values 0, 1, 2, 3, 4, 5, or 6.

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

**Phoenix Cardiovascular Scoring**

The Phoenix Cardiovascular score ranges from 0 to 6 points; 0, 1, or 2 points for each of systolic vasoactive medications, lactate, and MAP.

*Systemic Vasoactive Medications*

0 medications	0 points
1 medication	1 point
2 or more medications	2 points

*Lactate*

[0, 5)	0 points
[5, 11)	1 point
[11, Inf)	2 points

*MAP*

Age in [0, 1) months	[31, Inf) mmHg	0 points
	[17, 31) mmHg	1 point
	[0, 17) mmHg	2 points
Age in [1, 12) months	[39, Inf) mmHg	0 points
	[25, 39) mmHg	1 point
	[0, 25) mmHg	2 points
Age in [12, 24) months	[44, Inf) mmHg	0 points
	[31, 44) mmHg	1 point
	[0, 31) mmHg	2 points
Age in [24, 60) months	[45, Inf) mmHg	0 points
	[32, 45) mmHg	1 point

Age in [60, 144) months	[0, 32) mmHg	2 points
	[49, Inf) mmHg	0 points
	[36, 49) mmHg	1 point
Age in [144, 216] months	[0, 36) mmHg	2 points
	[52, Inf) mmHg	0 points
	[38, 52) mmHg	1 point
	[0, 38) mmHg	2 points

## References

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

## See Also

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

## Examples

```
# using the example sepsis data set
phoenix_cardiovascular(
  vasoactives = dobutamine + dopamine + epinephrine + milrinone + norepinephrine + vasopressin,
  lactate = lactate,
  age = age,
  map = dbp + (sbp - dbp)/3,
  data = sepsis
)

# example data set to get all the possible scores
DF <-
  expand.grid(vasos = c(NA, 0:6),
             lactate = c(NA, 3.2, 5, 7.8, 11, 14),
             age = c(NA, 0.4, 1, 3, 12, 18, 24, 45, 60, 61, 144, 145),
             map = c(NA, 16:52))
DF$card <- phoenix_cardiovascular(vasos, lactate, age, map, DF)
```

```

head(DF)

# what if lactate is unknown for all records? - set the value either in the
# data object or the arguement value to NA
DF2 <-
  expand.grid(vasos = c(NA, 0:6),
             age = c(NA, 0.4, 1, 3, 12, 18, 24, 45, 60, 61, 144, 145),
             map = c(NA, 16:52))
DF2$card <- phoenix_cardiovascular(vasos, lactate = NA, age, map, DF2)

DF3 <-
  expand.grid(vasos = c(NA, 0:6),
             lactate = NA,
             age = c(NA, 0.4, 1, 3, 12, 18, 24, 45, 60, 61, 144, 145),
             map = c(NA, 16:52))
DF3$card <- phoenix_cardiovascular(vasos, lactate, age, map, DF3)

identical(DF2$card, DF3$card)

```

---

phoenix\_coagulation    *Phoenix Coagulation Score*

---

## Description

Applies the Phoenix coagulation organ dysfunction scoring to a set of inputs.

## Usage

```

phoenix_coagulation(
  platelets = NA_real_,
  inr = NA_real_,
  d_dimer = NA_real_,
  fibrinogen = NA_real_,
  data = parent.frame(),
  ...
)

```

## Arguments

platelets	numeric vector for platelets counts in units of 1,000/uL (thousand per microliter)
inr	numeric vector for the international normalised ratio blood test
d_dimer	numeric vector for D-Dimer, units of mg/L FEU
fibrinogen	numeric vector units of mg/dL
data	a list, data.frame, or environment containing the input vectors
...	pass through

**Value**

a integer vector with values 0, 1, or 2

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

**Phoenix Coagulation Scoring**

1 point each for platelets < 100 K/micro liter, INR > 1.3, D-dimer > 2 mg/L FEU, and fibrinogen < 100 mg/dL, with a max total score of 2.

**References**

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

**See Also**

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

**Examples**

```
# using the example data set
phoenix_coagulation(
  platelets = platelets,
  inr = inr,
  d_dimer = d_dimer,
  fibrinogen = fibrinogen,
  data = sepsis
)

# build a data.frame with values for all possible combinations of values
# leading to all possible coagulation scores.
DF <-
  expand.grid(plts = c(NA, 20, 100, 150),
             inr = c(NA, 0.2, 1.3, 1.8),
             ddmr = c(NA, 1.7, 2.0, 2.8),
             fib = c(NA, 88, 100, 120))
```

```
DF$coag <- phoenix_coagulation(plts, inr, ddmr, fib, DF)
DF
```

---

phoenix\_endocrine      *Phoenix Endocrine Score*

---

## Description

Assess the Phoenix endocrine organ dysfunction score. This score is not part of the Phoenix score, only part of the Phoenix-8 score.

## Usage

```
phoenix_endocrine(glucose = NA_real_, data = parent.frame(), ...)
```

## Arguments

glucose	numeric vector; blood glucose measured in mg/dL
data	a list, data.frame, or environment containing the input vectors
...	pass through

## Value

a integer vector with values 0 or 1

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

## Phoenix Endocrine Scoring

The endocrine dysfunction score is based on blood glucose with one point for levels < 50 mg/dL or > 150 mg/dL.

## References

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

## See Also

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and

- phoenix\_endocrine,
- phoenix\_immunologic,
- phoenix\_renal,
- phoenix\_hepatic,

vignette('phoenix') for more details and examples.

## Examples

```
# using the example sepsis data set
endo_example <- sepsis[c("pid", "glucose")]
endo_example$score <- phoenix_endocrine(glucose, data = sepsis)
endo_example

# example data set to get all the possible endocrine scores
DF <- data.frame(glc = c(NA, 12, 50, 55, 100, 150, 178))
phoenix_endocrine(glucose = glc, data = DF)
```

---

phoenix_hepatic	<i>Phoenix Hepatic Score</i>
-----------------	------------------------------

---

## Description

Apply the Phoenix Hepatic scoring based on total bilirubin and ALT.

## Usage

```
phoenix_hepatic(
  bilirubin = NA_real_,
  alt = NA_real_,
  data = parent.frame(),
  ...
)
```

## Arguments

bilirubin	numeric vector; units of mg/dL
alt	alanine aminotransferase; a numeric vector; units of IU/L
data	a list, data.frame, or environment containing the input vectors
...	pass through

## Value

a integer vector with values 0 or 1

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

## Phoenix Hepatic Scoring

1 point for total bilirubin greater or equal to 4 mg/dL and/or ALT strictly greater than 102 IU/L.

## References

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

## See Also

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

## Examples

```
# using the example sepsis data set
hep_example <- sepsis[c("pid", "bilirubin", "alt")]
hep_example$score <- phoenix_hepatic(bilirubin, alt, sepsis)
hep_example

# example data set with all possible hepatic scores
DF <- expand.grid(bil = c(NA, 3.2, 4.0, 4.3), alt = c(NA, 99, 102, 106))
phoenix_hepatic(bilirubin = bil, alt = alt, data = DF)
```

---

phoenix\_immunologic    *Phoenix Immunologic Score*

---

## Description

Apply the Phoenix immunologic scoring based on ANC and ALC. This is only part of Phoenix-8 and not Phoenix.

## Usage

```
phoenix_immunologic(anc = NA_real_, alc = NA_real_, data = parent.frame(), ...)
```



## Arguments

<code>anc</code>	absolute neutrophil count; a numeric vector; units of 1,000 cells per cubic millimeter
<code>alc</code>	absolute lymphocyte count; a numeric vector; units of 1,000 cells per cubic millimeter
<code>data</code>	a list, data.frame, or environment containing the input vectors
<code>...</code>	pass through

## Value

a integer vector with values 0 or 1

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

## Phoenix Immunologic Scoring

1 point if ANC < 500 or ALC < 1000 cells per cubic millimeter.

## References

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

## See Also

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

## Examples

```
# using the example sepsis data set
immu_example <- sepsis[c("pid", "anc", "alc")]
immu_example$score <- phoenix_immunologic(anc, alc, sepsis)
immu_example

# using the example sepsis data set
hep_example <- sepsis[c("pid", "bilirubin", "alt")]
```

```

hep_example$score <- phoenix_hepatic(bilirubin, alt, sepsis)
hep_example

# example data set with all possible hepatic scores
DF <- expand.grid(anc = c(NA, 200, 500, 600), alc = c(NA, 500, 1000, 2000))
phoenix_immunologic(anc = anc, alc = alc, data = DF)

```

---

phoenix\_neurologic      *Phoenix Sepsis Neurological Score*

---

### Description

Assessment of neurologic dysfunction based on Glasgow Coma Scale (GCS) and pupil reactivity. This score is part of the diagnostic Phoenix Sepsis criteria and Phoenix 8 Sepsis criteria.

### Usage

```

phoenix_neurologic(
  gcs = NA_integer_,
  fixed_pupils = NA_real_,
  data = parent.frame(),
  ...
)

```

### Arguments

<code>gcs</code>	integer vector; total Glasgow Comma Score
<code>fixed_pupils</code>	integer vector; 1 = bilaterally fixed pupil, 0 = otherwise
<code>data</code>	a list, data.frame, or environment containing the input vectors
<code>...</code>	pass through

### Details

Missing values will map to a value of 0 as was done when developing the Phoenix criteria. Note that this is done on a input by input basis. That is, if pupil reactivity is missing but GCS (total) is 9, then the neurologic dysfunction score is 1.

GCS total is the sum of a score based on eyes, motor control, and verbal responsiveness.

Eye response:

1. no eye opening,
2. eye opening to pain,
3. eye opening to sound,
4. eyes open spontaneously.

Verbal response:

1. no verbal response,
2. incomprehensible sounds,
3. inappropriate words,
4. confused,
5. orientated

Motor response:

1. no motor response,
2. abnormal extension to pain,
3. abnormal flexion to pain,
4. withdrawal from pain,
5. localized pain,
6. obeys commands

### Value

an integer vector with values 0, 1, or 2. As with all Phoenix organ dysfunction scores, missing input values map to scores of zero.

### Phoenix Neurological Scoring

Bilaterally fixed pupil	2 points
Glasgow Coma Score (total) less or equal 10	1 point
Reactive pupils and GCS > 10	0 point

### References

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

### See Also

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

**Examples**

```
# using the example sepsis data set
phoenix_neurologic(
  gcs = gcs_total,
  fixed_pupils = as.integer(pupil == "both-fixed"),
  data = sepsis
)

# build an example data set with all possible neurologic scores
DF <- expand.grid(gcs = c(3:15, NA), pupils = c(0, 1, NA))
DF$target <- 0L
DF$target[DF$gcs <= 10] <- 1L
DF$target[DF$pupils == 1] <- 2L
DF$current <- phoenix_neurologic(gcs, pupils, DF)
stopifnot(identical(DF$target, DF$current))
DF
```

---

phoenix\_renal

*Phoenix Renal Score*


---

**Description**

Apply the Phoenix renal organ dysfunction score based on age adjusted creatinine levels.

**Usage**

```
phoenix_renal(
  creatinine = NA_real_,
  age = NA_real_,
  data = parent.frame(),
  ...
)
```

**Arguments**

creatinine	numeric vector; units of mg/dL
age	numeric vector age in months
data	a list, data.frame, or environment containing the input vectors
...	pass through

**Value**

a integer vector with values 0, 1, or 2

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

**Phoenix Renal Scoring**

Age in [0, 1) months	creatinine [0, 0.8) mg/dL	0 points
	creatinine [0.8, Inf) mg/dL	1 point
Age in [1, 12) months	creatinine in [0, 0.3) mg/dL	0 points
	creatinine in [0.3, Inf) mg/dL	1 point
Age in [12, 24) months	creatinine in [0, 0.4) mg/dL	0 points
	creatinine in [0.4, Inf) mg/dL	1 point
Age in [24, 60) months	creatinine in [0, 0.6) mg/dL	0 points
	creatinine in [0.6, Inf) mg/dL	1 point
Age in [60, 144) months	creatinine in [0, 0.7) mg/dL	0 points
	creatinine in [0.7, Inf) mg/dL	1 point
Age in [144, 216] months	creatinine in [0, 1.0) mg/dL	0 points
	creatinine in [1.0, Inf) mg/dL	1 point

**References**

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

**See Also**

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

**Examples**

```
# using the example sepsis data set
renal_example <- sepsis[c("creatinine", "age")]
renal_example$score <- phoenix_renal(creatinine, age, sepsis)
renal_example

# build an example data set with all possible neurologic scores
DF <- expand.grid(age = c(NA, 0.4, 1, 3, 12, 18, 24, 45, 60, 61, 144, 145),
                 creatinine = c(NA, seq(0.0, 1.1, by = 0.1)))
DF$card <- phoenix_renal(age = age, creatinine = creatinine, data = DF)

head(DF)
```

---

phoenix\_respiratory    *Phoenix Respiratory Score*

---

**Description**

Apply the Phoenix Respiratory Scoring rubric to a data set. The respiratory score is part of the diagnostic Phoenix Sepsis criteria and the diagnostic Phoenix 8 Sepsis criteria.

**Usage**

```
phoenix_respiratory(
  pf_ratio = NA_real_,
  sf_ratio = NA_real_,
  imv = NA_integer_,
  other_respiratory_support = NA_integer_,
  data = parent.frame(),
  ...
)
```

**Arguments**

<code>pf_ratio</code>	numeric vector for the PaO <sub>2</sub> /FiO <sub>2</sub> ratio; PaO <sub>2</sub> = arterial oxygen pressure; FiO <sub>2</sub> = fraction of inspired oxygen; PaO <sub>2</sub> is measured in mmHg and FiO <sub>2</sub> is from 0.21 (room air) to 1.00.
<code>sf_ratio</code>	numeric vector for the SpO <sub>2</sub> /FiO <sub>2</sub> ratio; SpO <sub>2</sub> = oxygen saturation, measured in a percent; ratio for 92% oxygen saturation on room air is 92/0.21 = 438.0952.
<code>imv</code>	invasive mechanical ventilation; numeric or integer vector, (0 = not intubated; 1 = intubated)
<code>other_respiratory_support</code>	other respiratory support; numeric or integer vector, (0 = no support; 1 = support)
<code>data</code>	a list, data.frame, or environment containing the input vectors
<code>...</code>	pass through

**Details**

`pf_ratio` is the ratio of partial pressure of oxygen in arterial blood (PaO2) to the fraction of inspiratory oxygen concentration (FiO2).

`sf_ratio` is a non-invasive surrogate for `pf_ratio` using pulse oximetry (SpO2) instead of invasive PaO2.

Important Note: when the Phoenix Sepsis criteria was developed there is a requirement that SpO2 < 97 in order for the `sf_ratio` to be valid. That assumption is not checked in this code and it is left to the end user to account for this when building the `sf_ratio` vector.

`imv` Invasive mechanical ventilation - integer vector where 0 = not intubated and 1 = intubated.

`other_respiratory_support` other respiratory support such as receiving oxygen, high-flow, non-invasive positive pressure, or imv.

**Value**

a integer vector with values 0, 1, 2, or 3.

As with all other Phoenix organ system scores, missing values in the data set will map to a score of zero - this is consistent with the development of the criteria.

**Phoenix Respiratory Scoring**

0 points

`pf_ratio` >= 400 and `sf_ratio` >= 292

1 point

(`pf_ratio` < 400 or `sf_ratio` < 292) and any respiratory support

2 points

(`pf_ratio` < 200 or `sf_ratio` < 200)

**References**

See reference details in [phoenix-package](#) or by calling `citation('phoenix')`.

**See Also**

- [phoenix](#) for generating the diagnostic Phoenix Sepsis score based on the four organ systems:
  - [phoenix\\_cardiovascular](#),
  - [phoenix\\_coagulation](#),
  - [phoenix\\_neurologic](#),
  - [phoenix\\_respiratory](#),
- [phoenix8](#) for generating the diagnostic Phoenix 8 Sepsis criteria based on the four organ systems noted above and
  - [phoenix\\_endocrine](#),
  - [phoenix\\_immunologic](#),
  - [phoenix\\_renal](#),
  - [phoenix\\_hepatic](#),

`vignette('phoenix')` for more details and examples.

**Examples**

```
# Using the provided example data set:
phoenix_respiratory(
  pf_ratio = pao2 / fio2,
  sf_ratio = spo2 / fio2,
  imv      = vent,
  other_respiratory_support = as.integer(fio2 > 0.21),
  data = sepsis
)

# A set of values that will get all possible respiratory scores:
DF <- expand.grid(
  pfr = c(NA, 500, 400, 350, 200, 187, 100, 56),
  sfr = c(NA, 300, 292, 254, 220, 177, 148, 76),
  vent = c(NA, 0, 1),
  o2 = c(NA, 0, 1)
)

phoenix_respiratory(
  pf_ratio = pfr,
  sf_ratio = sfr,
  imv = vent,
  other_respiratory_support = o2,
  data = DF
)
```

---

sepsis

*sepsis*


---

**Description**

A fully synthetic data set with variables need for examples and documentation of the Phoenix Sepsis Criteria.

**Usage**

```
sepsis
```

**Format**

a data.frame with 20 rows and 27 columns

[, 1]	pid	patient identification number
[, 2]	age	age in months
[, 3]	fio2	fraction of inspired oxygen
[, 4]	pao2	partial pressure of oxygen in arterial blood (mmHg)
[, 5]	spo2	pulse oximetry



[, 6]	vent	indicator for invasive mechanical ventilation
[, 7]	gcs_total	total Glasgow Coma Scale
[, 8]	pupil	character vector reporting if pupils are reactive or fixed.
[, 9]	platelets	platelets measured in 1,000 / microliter
[, 10]	inr	international normalized ratio
[, 11]	d_dimer	D-dimer; units of mg/L FEU
[, 12]	fibrinogen	units of mg/dL
[, 13]	dbp	diagnostic blood pressure (mmHg)
[, 14]	sbp	systolic blood pressure (mmHg)
[, 15]	lactate	units of mmol/L
[, 16]	dobutamine	indicator for receiving systemic dobutamine
[, 17]	dopamine	indicator for receiving systemic dopamine
[, 18]	epinephrine	indicator for receiving systemic epinephrine
[, 19]	milrinone	indicator for receiving systemic milrinone
[, 20]	norepinephrine	indicator for receiving systemic norepinephrine
[, 21]	vasopressin	indicator for receiving systemic vasopressin
[, 22]	glucose	units of mg/dL
[, 23]	anc	units of 1,000 cells per cubic millimeter
[, 24]	alc	units of 1,000 cells per cubic millimeter
[, 25]	creatinine	units of mg/dL
[, 26]	bilirubin	units of mg/dL
[, 27]	alt	units of IU/L

# Index

## \* datasets

sepsis, [24](#)

map, [2](#)

phoenix, [3](#), [5](#), [8](#), [11](#), [13](#), [14](#), [16](#), [17](#), [19](#), [21](#), [23](#)

phoenix8, [5](#), [6](#), [8](#), [11](#), [13](#), [14](#), [16](#), [17](#), [19](#), [21](#), [23](#)

phoenix\_cardiovascular, [5](#), [8](#), [9](#), [11](#), [13](#), [14](#),  
[16](#), [17](#), [19](#), [21](#), [23](#)

phoenix\_coagulation, [5](#), [8](#), [11](#), [12](#), [13](#), [14](#),  
[16](#), [17](#), [19](#), [21](#), [23](#)

phoenix\_endocrine, [5](#), [8](#), [11](#), [13](#), [14](#), [15–17](#),  
[19](#), [21](#), [23](#)

phoenix\_hepatic, [5](#), [8](#), [11](#), [13](#), [15](#), [15](#), [16](#), [17](#),  
[19](#), [21](#), [23](#)

phoenix\_immunologic, [5](#), [8](#), [11](#), [13](#), [15](#), [16](#),  
[16](#), [17](#), [19](#), [21](#), [23](#)

phoenix\_neurologic, [5](#), [8](#), [11](#), [13](#), [14](#), [16](#), [17](#),  
[18](#), [19](#), [21](#), [23](#)

phoenix\_renal, [5](#), [8](#), [11](#), [13](#), [15–17](#), [19](#), [20](#),  
[21](#), [23](#)

phoenix\_respiratory, [5](#), [8](#), [11](#), [13](#), [14](#), [16](#),  
[17](#), [19](#), [21](#), [22](#), [23](#)

sepsis, [24](#)