

# **Human CGRP ELISA Kit**

Enzyme Immunoassay for the quantification of CGRP human in serum and plasma samples

Catalog number: ARG80914

For research use only. Not for use in diagnostic procedures.

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#### INTRODUCTION

CALCA gene encodes the peptide hormones calcitonin, calcitonin gene-related peptide (CGRP) and katacalcin by tissue-specific alternative RNA splicing of the gene transcripts and cleavage of inactive precursor proteins. Calcitonin is involved in calcium regulation and acts to regulate phosphorus metabolism. Calcitonin gene-related peptide functions as a vasodilator and as an antimicrobial peptide while katacalcin is a calcium-lowering peptide. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Aug 2014]

CGRP induces vasodilation. It dilates a variety of vessels including the coronary, cerebral and systemic vasculature. Its abundance in the CNS also points toward a neurotransmitter or neuromodulator role. It also elevates platelet cAMP. [UniProt]

#### PRINCIPLE OF THE ASSAY

This is an Enzyme Immunoassay for the quantification CGRP in human serum and EDTA-plasma samples. (This kit is designed for serum and plasma samples but it may be used for other samples when the expression level is fall within the linear range.) This assay employs the competitive quantitative enzyme immunoassay technique. A secondary antibody has been pre-coated onto a microtiter plate. The secondary antibody can bind to the Fc fragment of the primary antibody which recognizes CGRP. The primary antibodies in the kit will be competitively bound by biotinylated-CGRP peptides and CGRP peptides in standards or targeted CGRP peptides in samples. The wells are washed and then incubated with Streptavidin-HRP reagent. The biotinylated peptide interacts with streptavidin-horseradish peroxidase to form a complex. After

washing away any unbound Streptavidin-HRP reagent, a substrate solution (TMB) is added to the wells and color develops in inverse-proportion to the amount of CGRP peptide present in the samples. The color development is stopped by the addition of STOP solution and the intensity of the color is measured at a wavelength of 450nm±2 nm. The concentration of CGRP peptide in the sample is then determined by comparing the O.D of samples to the standard curve.

### MATERIALS PROVIDED & STORAGE INFORMATION

Store the unopened kit at 2-8 °C. Use the kit before expiration date.

Component	Quantity	Storage information
Secondary antibody coated microplate	12 x 8 wells	4°C
20X Wash Buffer	50 ml	4°C
Primary antibody	1 vial	4°C
Biotinylated peptide	1 vial	4°C
Standard	1 vial	4°C
1000X Streptavidin-HRP conjugate	30 μΙ	4°C
TMB substrate	12 ml (Ready-to-use)	4°C (Protect from light)
STOP solution (2N HCl)	15 ml (Ready-to-use)	4°C
Positive Controls (Accept. Range dependent on lot#)	2 vials	4°C
Plate sealer	3 pieces	Room Temperature

## MATERIALS REQUIRED BUT NOT PROVIDED

- Microplate reader capable of measuring absorbance at 450nm
- Pipettes and pipette tips
- Deionized or distilled water
- Microplate shaker (300-400rpm)
- Aprotinin (enzyme inhibitor)
- C18 column (SEP- or SPE-COLUMN containing 200 mg of C18)
- Peptide extraction Binding Buffer (1% trifluoroacetic acid (TFA), HPLC grade is recommended)
- Peptide extraction Elution Buffer (60% acetonitrile, 1% TFA, and 39% distilled water, HPLC grade of acetonitrile and TFA are recommended)
- Automated microplate washer (optional)

#### TECHNICAL HINTS AND PRECAUTIONS

- Wear protective gloves, clothing, eye, and face protection especially while handling blood or body fluid samples.
- Store the kit at 4°C at all times.
- Before opening any Eppendorf tubes for reconstitution, briefly centrifuge at ~3,000rpm for 5 seconds to ensure that all the lyophilized material is at the bottom of the tube.
- If crystals are observed in the 20X Wash buffer, warm to RT or 37°C until the crystals are completely dissolved.
- Ensure complete reconstitution and dilution of reagents prior to use.
- It is highly recommended that all solutions be used as soon as possible after reconstitution.

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- Unused microplate strips should be placed back in the foil pouch with a desiccant and stored at 4°C. Do not allow moisture to enter the wells.
- All reagents should be mixed by gentle inversion or swirling prior to use.
  Do not induce foaming.
- Before using the kit, spin tubes and bring down all components to the bottom of tubes.
- Each time a new tip is used, make sure the tip is secure and free of air bubbles. For better intra-assay variation, aspirate and expel a reagent or sample back into the container a few times prior to loading.
- Avoid submerging the whole tip into reagents because droplets can accumulate at the end of the tip causing an excess of reagent to be loaded into the well. This can lead to poor results.
- For optimal results, an orbital plate shaker capable of 300-400 rpm is recommended for all incubations.
- It is highly recommended that the standards, samples and controls be assayed in duplicates.

### SAMPLE COLLECTION & STORAGE INFORMATION

The sample collection and storage conditions listed below are intended as general guidelines. Sample stability has not been evaluated. This kit is designed for serum and plasma samples but it may be used for other samples when the expression level is fall within the linear range.

<u>Serum</u>- Use a serum separator tube (SST) and allow samples to clot for 30 minutes before centrifugation for 15 minutes at 1600 x g at 4°C. Collect serum and assay immediately or aliquot and store samples at  $\leq$  -80 °C up to 1 month. Avoid repeated freeze-thaw cycles.

<u>Plasma</u> - Collect plasma using EDTA as an anticoagulant. Centrifuge for 15 minutes at 4°C at 1000 x g within 30 minutes of collection. Assay immediately or aliquot and store samples at  $\leq$  -80 °C up to 1 month. Avoid repeated freezethaw cycles.

**Note:** We recommended add Aprotinin (enzyme inhibitor) for **ALL** sample collection to prevent sample degradation. 0.6 TIU or 100  $\mu$ l of Aprotinin per mL of sample solution.

# Peptide extraction for Serum/plasma-

- It is recommended samples can through peptide extraction protocol as below.
- 2. Mix an equal amount of Binding Buffer (1% trifluoroacetic acid (TFA), HPLC grade is recommended) with the plasma or serum samples and vortex the mixture. (E.g. 1ml of plasma/serum + 1 ml of Binding Buffer). Centrifuge at 6,000-17,000 × g for 20 minutes at 4°C. Collect the supernatant.
- 3. Slowly equilibrate a C18 column (SEP- or SPE-COLUMN containing 200 mg of C18) by washing the C18 column with 1 ml Elution Buffer (60% acetonitrile, 1% TFA, and 39% distilled water, HPLC grade of acetonitrile

- and TFA are recommended).
- **4.** Wash the C18 column with 3 ml of Binding Buffer three times.
- **5.** Load the sample/Binding Buffer mixture solution from step 2 into the washed C18 column from step 4.
- **6.** Wash the column slowly with Binding Buffer (3 ml, twice) and discard the wash.
- 7. Elute the peptide slowly with Elution Buffer (3 ml, once) and collect eluant into a polystyrene tube.
  - Note: From steps 5-7, no pressure should be applied to the column.
- **8.** Evaporate eluant to dryness in a centrifugal concentrator or by a suitable substitute method to dryness the eluant. (Freeze-dry the resulting water/TFA solution to dryness)
- 9. Keep the dried extract at-20°C and perform the assay as soon as possible.
- **10.** Use 1x wash buffer in the kit to reconstitute the dried extract. If the peptide value does not fall within the range of detection, dilute or concentrate the sample accordingly.

Note: For normal subject extracted from 1 ml original plasma/serum, use  $125 \,\mu l\,1x$  Wash buffer to reconstitute the dried extract. Aliquot  $50 \,\mu l$  into two designated assay wells ( $25 \,\mu l$  is left over). The concentration factor in this case is 8. (1 ml /  $125 \, ul = 8$ ). The original plasma peptide concentration level is 1/8 of the level of final extracted plasma.

e.g. If the level of the final extracted plasma is 100 pg/ml, then the total level of peptide in the original plasma = (100 pg/ml) / 8 = 12.5 pg/ml. After performing assay, if the peptide value exceeds or does not fall in the range of detection, dilute or concentrate the samples accordingly.

#### REAGENT PREPARATION

- **1X Wash Buffer**: Dilute 20X Wash Buffer into distilled water to yield 1X Wash buffer. Store the diluted 1X Wash Buffer at 4°C. If crystals appear in 20X Wash Buffer, warm the buffer in 37°C water bath for 30 minutes or until crystals disappear. Mix well before use.
- Primary antibody: Reconstitute the Primary antibody vial with 5 ml of 1X Wash Buffer. Allow it to sit for 5 minutes to completely dissolve, mix well and keep rehydrated solution at 4°C before use. Store the reconstituted antibody at 4°C up to a week. For long-term storage, aliquot & store at -20°C for up to 3 months. Avoid repeated freeze-thaw cycles. It is recommended that antibody should only be frozen-thawed once.
- Biotinylated peptide: Reconstitute the Biotinylated peptide vial with 5 ml of 1X Wash Buffer. Allow it to sit for 5 minutes to completely dissolve, mix well and keep rehydrated solution at 4°C before use. Store the reconstituted peptide at 4°C up to a week. For long-term storage, aliquot & store at -20°C for up to 3 months. Avoid repeated freeze-thaw cycles. It is recommended that peptide should only be frozen-thawed once.
- Positive control: Centrifuge and reconstitute the Positive control vial with 200 μl of 1X Wash Buffer. Allow it to sit for 5 minutes to completely dissolve, mix well and keep rehydrated solution at 4°C before use. Store the reconstituted control at 4°C up to a week. For long-term storage, aliquot & store at -20°C for up to 3 months. Avoid repeated freeze-thaw cycles. It is recommended that control should only be frozen-thawed once. (acceptable range dependent on lot#)

- 1X Streptavidin-HRP conjugate: 1X Streptavidin-HRP conjugate working solution should be prepared freshly before use. Centrifuge 1000X Streptavidin-HRP conjugate briefly and add 12µl of Streptavidin-HRP to 12ml 1X wash buffer to make a 1X Streptavidin-HRP working solution. Vortex thoroughly.
- Standard peptide: Centrifuge and reconstitute the standard with 1 ml of 1X Wash buffer and vortex. The concentration of this stock solution is 1000 ng/ml. Allow the solution to sit for at least 10 minutes at room temperature to completely dissolve. Store the reconstituted standard at 4°C up to a week. For long-term storage, aliquot & store at -20°C for up to 3 months. Avoid repeated freeze-thaw cycles. It is recommended that standard should only be frozen-thawed once. Dilute peptide standard solutions with 1X Wash buffer to 100 ng/ml, 10 ng/ml, 1 ng/ml, 0.1 ng/ml, 0.01 ng/ml as follows:

The example of the dilution of standards

Standard No.	Standard Conc. (ng/ml)	1X Wash Buffer (μl)	Standard (µl)
Stock	1000	-	-
S1	100	900	100 μl of Stock
S2	10	900	100 μl of S1
S3	1	900	100 μl of S2
S4	0.1	900	100 μl of S3
S5	0.01	900	100 μl of S4
SO (Total binding)	0	900	0

### **ASSAY PROCEDURE**

All materials should be equilibrated to room temperature (RT, 20-23°C) 30 minutes before opening and starting the assay. Standards, samples and blank should be assayed in duplicates.

- 1. Remove excess microtiter strips from the plate frame, return them to the foil pouch containing the desiccant pack, and reseal it.
- 2. Add **50 μl of 1X Wash Buffer** as Total Binding (S0, zero standard). Two empty wells should be left as blank.
- 3. Add  $50 \,\mu$ l of prediluted peptide standards (add from S5 to S1),  $50 \,\mu$ l positive controls or  $50 \,\mu$ l samples into corresponding wells. It is advisable to assay each condition in duplicates.
- 4. Add 25 μl of primary antibody into each well except the Blank wells.
- 5. Add  $25 \,\mu l$  of Biotinylated peptide into each well except the Blank wells. It is not recommended to use a multi-channel pipette to load the primary antibody and biotinylated peptide.
- Seal the microtiter plate with plate sealer. Incubate for 2 hours at RT (20-23°C). Orbital shaking on a microplate shaker at 300-400 rpm is recommended.
- 7. Prepare 1X Streptavidin-HRP conjugate working solution: Mix and centrifuge Streptavidin-HRP concentrate vial (3,000-5,000 rpm for 5 seconds) before use. Pipette 12  $\mu$ l of Streptavidin-HRP concentrate into 12 ml of 1X Wash Buffer to make a Streptavidin-HRP working solution. Vortex thoroughly. Prepare freshly.
- 8. Remove sealer from plate.
- 9. Aspirate each well and wash, repeating the process 3 times for a total 4

washes. Wash by filling each well with  $1\times$  Wash Buffer (350  $\mu$ I) using a squirt bottle, manifold dispenser, or autowasher. Complete removal of liquid at each is essential to good performance. After the last wash, remove any remaining Wash Buffer by aspirating, decanting or blotting against clean paper towels.

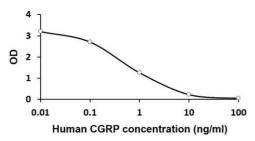
- 10. Add **100 μl of diluted (1X) Streptavidin-HRP** working solution into **each well**.
- 11. Reseal the plate with sealer. Incubate for **1 hour at RT**. Orbital shaking on a microplate shaker at 300-400 rpm is recommended.
- 12. Remove sealer from plate. Wash as according to step 9.
- 13. Add 100 µl of TMB substrate solution into each well.
- 14. Reseal the plate with sealer. Incubate for **1** hour at RT in dark. Orbital shaking on a microplate shaker at 300-400 rpm is recommended.
- 15. Remove sealer from plate. (DO NOT wash or discard the contents of the wells)
- 16. Add 100  $\mu$ l of STOP solution (2N HCl) into each wells to stop the reaction. Gently tap the plate to ensure thorough mixing. The color of the solution should change from blue to yellow.
- 17. Read the OD with a microplate reader at **450 nm** immediately. It is recommended that the wells be read <u>within 20 minutes</u> after adding the Stop Solution.

### **CALCULATION OF RESULTS**

- 1. Calculate the average absorbance values for each set of standards, controls and patient samples.
- 2. Using log-log, semi-log or linear graph paper, construct a standard curve by plotting the mean absorbance obtained from each standard against its concentration with absorbance value on the vertical (Y) axis and concentration on the horizontal (X) axis.
- 3. Using the mean absorbance value for each sample determine the corresponding concentration from the standard curve.
- 4. Automated method: The results in the IFU have been calculated automatically using a 4 PL (4 Parameter Logistics) curve fit. 4 Parameter Logistics is the preferred method. Other data reduction functions may give slightly different results.
- 5. If samples have been diluted prior to the assay, the measured concentration must be multiplied by their respective dilution factors.

### **EXAMPLE OF TYPICAL STANDARD CURVE**

The following data is for demonstration only and cannot be used in place of data generations at the time of assay.



# **QUALITY ASSURANCE**

# Sensitivity

The standard of CGRP peptide ranged from 0-100 ng/ml.

The mean MDD was 0.28 ng/ml.

# **Linear Range**

0.28 - 3.38 ng/ml

### Precision:

Intra-assay: < 10%

Inter-assay: < 15%

# **Cross Reactivity**

The cross reactivity ratio of the tested peptide as the table:

Peptide	Cross Reactivity (%)
CGRP (Human)	100
CGRP (8-37) (Human)	100
CGRP (Chicken)	100
CGRP (Rat)	100
CGRP II (Human)	100
CGRP II (Rat)	100
Calcitonin (Human)	0
CAP (Rat)	0
Amylin (Human)	0
Amylin Amide (Human)	0
Amylin (Rat)	0
Substance P	0

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