

Immunoglobulin G (IgG) is antibody molecule. Each IgG is composed of four peptide chains — two heavy chains γ and two light chains. IgG antibodies are large molecules of about 150 kDa composed of four peptide chains. It contains two identical class γ heavy chains of about 50 kDa and two identical light chains of about 25 kDa, thus a tetrameric quaternary structure. The two heavy chains are linked to each other and to a light chain each by disulfide bonds. The resulting tetramer has two identical halves, which together form the Y-like shape. Each end of the fork contains an identical antigen binding site. The Fc regions of IgGs bear a highly conserved N-glycosylation site. The N-glycans attached to this site are predominantly core-fucosylated diantennary structures of the complex type. In addition, small amounts of these N-glycans also bear bisecting GlcNAc and α -2, 6-linked sialic acid residues [1]. IgG antibodies are involved in predominantly the secondary immune response. The presence of specific IgG, in general, corresponds to maturation of the antibody response [2]. IgG can bind to many kinds of pathogens, for example viruses, bacteria, and fungi, and protects the body against them by agglutination and immobilization, complement activation (classical pathway), opsonization for phagocytosis, and neutralization of their toxins.

There are four IgG subclasses (IgG1, 2, 3, and 4), named in order of their abundance in serum and IgG2 is the second most abundant. The relative ability of different IgG subclasses to fix complement may explain why some anti-donor antibody responses do harm a graft after organ transplantation. ^[3] In a Canine model of autoantibody mediated anemia using IgG isotype switch variants of an anti-erythrocytes autoantibody, it was found that Canine IgG2a was superior to IgG1 in activating complement. Moreover, it was found that the IgG2a isotype was able to interact very efficiently with Fc gamma R. As a result, 20 times higher doses of IgG1, in relationship to IgG2a autoantibodies, were required to induce autoantibody mediated pathology. ^[4]

References

- 1. Stadlmann J, et al. (2008). Proteomics 8 (14): 2858.
- 2. Meulenbroek, A.J. et al. (1996).
- 3. Gao ZH et al. (2004). Liver Transplantation 10 (8): 1055–1059.
- 4. Azeredo da Silveira S, et al. (2002). *J Exp Med* **195**(6): 665–72.

PRINCIPLE OF THE ASSAY

This is a shorter ELISA assay that reduces time to 50% compared to the conventional method, and the entire assay only takes 3 hours. This assay employs the quantitative sandwich enzyme immunoassay technique and uses biotin-streptavidin chemistry to improve the performance and the sensitivity of the assays. The sensitivity is 125-folds higher than the conventional method. An antibody specific for Canine IgG2 has been pre-coated onto a microplate. Standards and samples are pipetted into the wells and any IgG2 present is bound by the immobilized antibody. After washing away any unbound substances, a detection antibody specific for Canine IgG2 is added to the wells. Following wash to remove any unbound antibody reagent, a detection reagent is added. After intensive wash a substrate solution is added to the wells and color develops in proportion to the amount of IgG2 bound in the initial step. The color development is stopped and the intensity of the color is measured.

This package insert must be read in its entirety before using this product.

Storage

Store the kit at 4°C. The kit can be used in 6 months.



MATERIALS PROVIDED

Description	Quantity	Description	Quantity	Description	Quantity
Antibody Precoated Plate	1	20 x PBS	1	Substrate Solution	1
Detection Antibody	1	20 x Assay Buffer	1	Stop Solution	1
HRP Conjugate	1	96-well plate sheet	1	DataSheet/Manual	1
Standard	3				

Bring all reagents to room temperature before use.

Reagent Preparations

Canine IgG2 Detection Antibody (1 vial) – The lyophilized Detection Antibody should be stored at 4° C or -20°C for up to 6 months, if not used immediately. Centrifuge for 1 min at 6000 x g to bring down the material prior to open the vial. The vial contains sufficient Detection Antibody for a 96-well plate. Add 200 μ L of 1 x PBS to the antibody vial and vortex 15 sec and allow it to sit 5 min. Take 200 μ L of the detection antibody to 10.5 mL of 1 x PBS to make working dilution of Detection Antibody if the entire 96-well plate is used. If the partial antibody is used store the rest at -20°C until use.

Canine IgG2 Standard (3 vials) – Each of the lyophilized Canine IgG2 Standard vial contains the standard sufficient for generating a calibration curve. The unreconstituted standard can be stored at 4°C or -20°C for up to 6 months if not used immediately. Centrifuge for 1 min at 6000 x g to bring down the material prior to open the tube. Add 500 μ L of 1 x Assay Buffer to a standard vial to make the high standard concentration of 20 ng /ml, vortex for 15 sec and allow it to sit for 5 min. A seven-point standard curve is generated using 2-fold serial dilutions in the Assay Buffer, vortex 20 sec for each of dilution step.

HRP Conjugate (55 μ L) – Centrifuge for 1 min at 6000 x g to bring down the material prior to open the vial. The vial contains 55 μ L HRP Conjugate sufficient for one 96-well plate. If the volume is less than 55 μ L, add sterile 1 x PBS to reach 55 μ L and vortex briefly. Make 1:200 dilution in 1 x PBS. If the entire 96-well plate is used, add 53 μ L of HRP Conjugate to 10.5 mL of the PBS to make working dilution of HRP Conjugate prior to the assay. The rest of undiluted HRP Conjugate can be stored at 2 - 8°C for up to 6 months. DO NOT FREEZE

20 x PBS, pH 7.3, 25 mL- Dilute to 1 x PBS with deionized distilled water and mix well prior to use.

20 x Assay Buffer, 20 mL- Dilute to 1 x Assay Buffer with 1 x PBS prior to use.

Substrate Solution, 10.5 mL.

Stop Solution, 5.5 mL.



Assay Procedure

- 1. All procedures are conducted at room temperature (20-25 °C) and ensure **equal pipetting/dispensing** at each step and remove air bubbles in the wells for all steps.
- 2. Lift the plate cover and cover the unused wells or reseal the unused strips in the aluminum bag with desiccant at 4 °C. Vortex the standards and samples for 10 sec before applying to the plate. Add 100 μL of sample or standard per well and use duplicate wells for each standard or sample. Cover the 96-well plate and incubate for 2 hours. Attention: MUST vortex standards and samples for 10 sec before pipetting to the wells!
- 3. Aspirate each well and wash with 300 µL of 1 x Assay Buffer for two times. Wash by filling each well with 1 x Assay Buffer using a multi-channel pipette, manifold dispenser or autowasher. Complete removal of liquid at each step is essential for good performance. After the last wash, remove any remaining Assay Buffer by aspirating or by inverting the plate and blotting it against clean paper towels.
- 4. Add 100 μL of the **working dilution of Detection Antibody** to each well. Cover the plate and incubate for 1 hour.
- 5. Repeat the aspiration/wash as in step 3.
- 6. Add 100 μL of the **working dilution of HRP Conjugate** to each well. Cover the plate and incubate for 20 minutes. Avoid placing the plate in direct light.
- 7. Repeat the aspiration/wash as in step 3 but wash 4 times instead.
- 8. Add 100 μL of **Substrate Solution** to each well and observe the color development every 1-2 mins. Incubate for up to **30 minutes** (**depending on signal. Stop** the reaction when it turns to dark blue in the highest standard wells). Over-incubation of the substrate will result in overflow of high standard and thus should be avoided. Avoid placing the plate in direct light.
- 9. When it gets to dark blue in the highest concentration of standard wells, add 50 μL of **Stop Solution** to each well to stop the reaction. Gently tap the plate to ensure thorough mixing.
- 10. Determine the optical density of each well immediately, using a microplate reader set to 450 nm. If wavelength correction is available, set to 540 nm or 570 nm. If wavelength correction is not available, subtract readings at 540 nm or 570 nm from the readings at 450 nm. This subtraction will correct for optical imperfections in the plate. Readings made directly at 450 nm without correction may be higher and less accurate.

Sample dilution: If high density is expected samples should be diluted with one volume of 1 x Assay Buffer and **vortex for 1 min** prior to assay. If the OD value still exceeds the upper limit of the standard curve, further dilution is recommended till it falls in the detection range and the dilution factor must be used for calculation of the concentration.

Precaution and Technical Notes

- 1. It is critical to follow the procedure step by step otherwise appropriate color development may not occur as expected and make sure no air bubbles in wells before adding reagents.
- 2. A standard curve should be generated for each set of samples assayed. Thorough mixing of standards at each of dilution steps is critical to acquire a normal standard curve and **vortex again (10 sec) before pipetting to the 96-well plate**.
- 3. HRP Conjugate contains enzyme, DO NOT mass up with Detection Antibody.
- 4. The Stop Solution is an acid solution, handle with caution.
- 5. This kit should not be used beyond the expiration date on the label.
- 6. A thorough and consistent wash technique is essential for proper assay performance.
- 7. Use a fresh reagent reservoir and pipette tips for each step.
- 8. It is recommended that all standards and samples be assayed in duplicate.
- 9. Avoid microbial contamination of reagents and buffers. This may interfere with the performance of the assay.



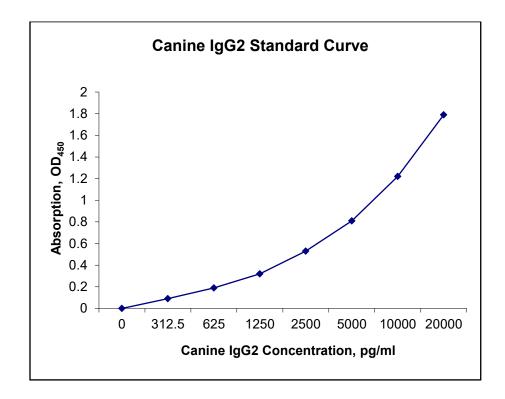
Calculation of Results

Average the duplicate readings for each standard, control, and sample and subtract the average zero (blank) standard optical density.

Create a standard curve by reducing the data using computer software capable of generating a four-parameter logistic (4-PL) curve-fit. As an alternative, construct a standard curve by plotting the mean absorbance for each standard on the y-axis against the concentration on the x-axis and draw a best fit curve through the points on the graph. The data may be linearized by plotting the log of the IgG2 concentrations versus the log of the O.D. and the best fit line can be determined by regression analysis. This procedure will produce an adequate but less precise fit of the data. If samples have been diluted, the concentration read from the standard curve must be multiplied by the dilution factor.

The Standard Curve

The graph below represents typical data generated when using this Canine IgG2 ELISA Kit. The standard curve was calculated using a computer generated 4-PL curve-fit. For this case, a Bio-Rad iMarkTM Microplate Reader and a Microplate Manager 6 Software were used to generate this curve. The correlation coefficient (r²) is 0.999-1.000.





Specificity

The following recombinant Canine proteins prepared at 10 ng/ml were tested and exhibited no cross-reactivity or interference.

BMP1, IgA, IgE, IL-1 β , IFN γ , TGF β 1, TLR3, TNF- α , VEGF.

Calibration

This kit is calibrated against canine IgG2.

Detection Range

0.31-20 ng/ml

Assay Sensitivity

60 pg/ml

Assay Precision

Intra-Assay %CV: 5; Inter-Assay %CV: 9

Related products

Canine IgG2 Standard
Canine IgG2 detection antibody

DECLARATION

THIS REAGENT IS FOR IN VITRO LABORATORY TESTING AND RESEARCH USE ONLY. DO NOT USE IT FOR CLINICAL DIAGNOSTICS. DO NOT USE OR INJECT IT IN HUMANS AND ANIMALS.

FOR LABORATORY RESEARCH USE ONLY NOT FOR USE IN HUMANS AND ANIMALS



Troubleshooting Guide

Troubleshootin		
Problem	Possible causes	Solution
Poor standard curve	 Inaccurate pipetting Insufficient vortexing OD₄₅₀ too high for the high standard point Air bubbles in wells. 	 Check pipettes and ensure equal dispensing. Vortex 30 sec for each of standard dilution steps and vortex again (10 sec) before pipetting to the 96-well plate. Reduce substrate incubation time Remove air bubbles in wells by pipette tip.
Low signal	 Improper preparation of reagents and storage Too brief incubation times Inadequate reagent volume or improper dilution Standard defect 	 Briefly spin down vials before opening. Reconstitute the powder thoroughly. Proper storage of plate and strip and detection antibody after first usage as shown in the datasheet. Ensure sufficient incubation time including substrate incubation. Increase sample incubation to 2 hours. Change a Standard vial.
Overflow in the standards	 Substrate incubation too long Air bubbles in wells 	 Observe the color development every 1-2 mins and reduce substrate incubation time. Remove air bubbles in wells Stop the reaction by adding 50 µl of Stop Solution when it turns to dark blue in the highest concentration of standard wells.
Large CV	 Inaccurate pipetting and mixing Improper standard/sample dilutions. Air bubbles in wells. 	 Check pipettes and ensure accurate pipetting and thorough mixing and equal dispensing. Use the correct dilution buffers Remove air bubbles in wells by pipette tip.
High background	 Reagent reservoir issue Plate is insufficiently washed and air bubbles in wells. Contaminated wash buffer Pipet tip contaminated 	 Use a new reagent reservoir for Substrate Solution. Increase wash to 4 times before adding substrate and ensure plate washer functions normally. Remove air bubbles in wells by pipette tip. Make fresh wash buffer and wash thoroughly. Use new pipette tips for blank wells.
No signal detected	 The procedure was misconducted. Failures of spin down the contents in Detection Antibody and Standards. Failure of Substrate or HRP Samples overdiluted 	 Ensure the step-by-step protocol. Spin vials of Detection antibody and Standard to complete recover the content. Mix 100 μl of Substrate with 0.5 μl HRP and deep blue color should develop in 2 min. Try a new standard vial and use positive control. Try not dilute samples
Low sensitivity	 Improper dilutions of standards Improper storage of the ELISA kit 	 Ensure accurate and thorough dilutions of standards at each step. Store detection antibody at -20°C after reconstitution, others at 4°C. Keep substrate solution protected from light.