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Datasheet for ABIN454984 Dopamine Receptor d1 ELISA Kit



Overview

Quantity:	96 tests
Target:	Dopamine Receptor d1 (DRD1)
Reactivity:	Human
Method Type:	Sandwich ELISA
Detection Range:	0.156-10 ng/mL
Minimum Detection Limit:	0.156 ng/mL
Application:	ELISA
Product Details	
Purpose:	This immunoassay kit allows for the specific measurement of human Dopamine D1 receptor
	(D1R) concentrations in cell culture supernates, serum, and plasma.
Sample Type:	Cell Culture Supernatant, Serum, Plasma
Analytical Method:	Quantitative
Detection Method:	Colorimetric
Specificity:	This assay recognizes recombinant and natural human D1R.
Cross-Reactivity (Details):	No significant cross-reactivity or interference was observed.
Characteristics:	Homo sapiens,Human,D(1A) dopamine receptor,Dopamine D1 receptor,DRD1
Components:	Reagent (Quantity): Assay plate (1), Standard (2), Sample Diluent (1x20ml), Assay Diluent A
	(1x10ml), Assay Diluent B (1x10ml), Detection Reagent A 1×120µl 2 Detection Reagent B
	1×120µl Wash Buffer(25 x concentrate) (1x30ml), Substrate (1x10ml), Stop Solution (1x10ml)

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Target Details	
Target:	Dopamine Receptor d1 (DRD1)
Alternative Name:	DRD1 (DRD1 Products)
Background:	Dopamine is an endogenous catecholamine that influences many cellular activities, including behavior, hormone synthesis and release, blood pressure and intracellular ion transport. A family of at least 5 Dopamine Receptors (DR) genes, D1-D5, have been identified based upon the amino acid identity, pharmacological specificity and effector responses. DR have been classified into either the D1-like (D1, D1B, and D5) or D2-like (D2, D3, and D4). The two isoforms of D2R, D2 long (D2L) and short D2S), are encoded by splice variants of a single gene and differ only by the presence of an additional 29 AA in the intracellular domain 3 of the D2 long form. It may play a role in the coupling of the receptor to G-proteins. All members of this family have similar structure and contain 7 putative transmembrane domains. A given cell or tissue may express more than one DR. Specific radioligands do not exist that can differentiate between these DR.
Pathways:	cAMP Metabolic Process, Inositol Metabolic Process, Protein targeting to Nucleus, Feeding Behaviour, Smooth Muscle Cell Migration, Regulation of long-term Neuronal Synaptic Plasticity

Application Details

Sample Volume:	100 μL
Plate:	Pre-coated
Protocol:	This assay employs the quantitative sandwich enzyme immunoassay technique. A monoclonal antibody specific for D1R has been pre-coated onto a microplate. Standards and samples are pipetted into the wells and any D1R present is bound by the immobilized antibody. An enzyme-linked monoclonal antibody specific for D1R is added to the wells. Following a wash to remove any unbound antibody-enzyme reagent, a substrate solution is added to the wells and color develops in proportion to the amount of D1R bound in the initial step. The color development is stopped and the intensity of the color is measured.
Reagent Preparation:	Bring all reagents to room temperature before use. Wash Buffer - If crystals have formed in the concentrate, warm to room temperature and mix gently until the crystals have completely dissolved. Dilute 20 mL of Wash Buffer Concentrate into deionized or distilled water to prepare 500 mL of Wash Buffer. Standard - Reconstitute the Standard with 1.0 mL of Sample Diluent. This reconstitution produces a stock solution of 10,000 pg/mL. Allow the standard to sit for a minimum of 15 minutes with gentle agitation prior to making serial dilutions. The undiluted standard serves as the high standard (10,000 pg/mL). The Sample Diluent serves as the zero

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Application Details	
	standard (0 pg/mL). Detection Reagent A and B - Dilute to the working concentration specified on the vial label using Assay Diluent A and B (1:100), respectively.
Sample Collection:	Cell culture supernates - Remove particulates by centrifugation and assay immediately or aliquot and store samples at \leq -20 °C. Avoid repeated freeze-thaw cycles. Serum - Use a serum separator tube (SST) and allow samples to clot for 30 minutes before centrifugation for 15 minutes at approximately 1000 x g. Remove serum and assay immediately or aliquot and store samples at -20 °C. Plasma - Collect plasma using EDTA or heparin as an anticoagulant. Centrifuge samples for 15 minutes at 1000 x g at 2 - 8 °C within 30 minutes of collection. Store samples at \leq -20 °C. Avoid repeated freeze-thaw cycles. Note: Citrate plasma has not been validated for use in this assay.
Assay Procedure:	 Allow all reagents to reach room temperature. Arrange and label required number of strips. 1. Prepare all reagents, working standards and samples as directed in the previous sections. 3 2. Add 100 uL of Standard, Control, or sample per well. Cover with the adhesive strip. Incubate for 2 hours at 37 °C. 3. Remove the liquid of each well, don't wash. 4. Add 100 uL of Detection Reagent A to each well. Incubate for 1 hour at 37°C. Detection Reagent A may appear cloudy. Warm to room temperature and mix gently until solution appears uniform. 5. Aspirate each well and wash, repeating the process three times for a total of three washes. Wash by filling each well with Wash Buffer (350 uL) using a squirt bottle, multi-channel pipette, manifold dispenser or autowasher. Complete removal of liquid at each step is essential to good performance. After the last wash, remove any remaining Wash Buffer by aspirating or decanting. Invert the plate and blot it against clean paper towels. 6. Add 100 uL of Detection Reagent B to each well. Incubate for 30 minutes at room temperature. Protect from light. 9. Add 50 uL of Stop Solution to each well. If color change does not appear uniform, gently tap the plate to ensure thorough mixing. 10. Determine the optical density of each well within 30 minutes, using a microplate reader set to 450 nm. Important Note: The wash procedure is critical. Insufficient washing will result in poor precision and falsely elevated absorbance readings.

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	2. It is recommended that no more than 32 wells be used for each assay run if manual pipetting
	is used since pipetting of all standards, specimens and controls should be completed within 5
	minutes. A full plate of 96 wells may be used if automated pipetting is available.
	3. Duplication of all standards and specimens, although not required, is recommended.
	4. When mixing or reconstituting protein solutions, always avoid foaming.
	5. To avoid cross-contamination, change pipette tips between additions of each standard level,
	between sample additions, and between reagent additions. Also, use separate reservoirs for
	each reagent.
	6. To ensure accurate results, proper adhesion of plate sealers during incubation steps is
	necessary.
Calculation of Results:	Average the duplicate readings for each standard, control, and sample and subtract the average
	zero 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 4 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 D1R 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.
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