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Does Anesthetic Additivity Imply a Similar Molecular Mechanism of Anesthetic Action at *N*-Methyl-d-Aspartate Receptors?

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Abstract

Background—Isoflurane and carbon dioxide (CO₂) negatively modulate N-methyl-D-aspartate (NMDA) receptors, but via different mechanisms. Isoflurane is a competitive antagonist at the NMDA receptor glycine binding site, whereas CO₂ inhibits NMDA receptor current through extracellular acidification. Isoflurane and CO₂ exhibit additive minimum alveolar concentration effects in rats, but we hypothesized that they would not additively inhibit NMDA receptor currents *in vitro* because they act at different molecular sites.

Methods—NMDA receptors were expressed in frog oocytes and studied using two-electrode voltage clamp techniques. A glycine concentration-response for NMDA was measured in the presence and absence of CO_2 . Concentration-response curves for isoflurane, H^+ , CO_2 , and ketamine as a function of NMDA inhibition were measured, and a Hill equation was used to calculate the EC_{50} for each compound.

Results—Binary drug combinations containing $\frac{1}{2}$ EC₅₀ were additive if NMDA current inhibition was not statistically different from 50%. The $\frac{1}{2}$ EC₅₀ binary drug combinations decreased the percent baseline NMDA receptor current as follows (mean±SD, n=5–6 oocytes each): CO₂+H⁺ (51±5%), CO₂+isoflurane (54±5%), H⁺+isoflurane (51±3%), CO₂+ketamine (67±8%), H⁺+ketamine (64±2%).

Conclusions—In contrast to our hypothesis, NMDA receptor inhibition by CO_2 and isoflurane is additive. Possibly, CO_2 acidification modulates a pH-sensitive loop on the NMDA receptor that in turn alters glycine binding affinity on the GluN1 subunit. However, ketamine plus either CO_2 or H⁺ synergistically inhibits NMDA receptor currents. Drugs acting via different mechanisms can thus exhibit additive or synergistic receptor effects. Additivity may not robustly indicate commonality between molecular anesthetic mechanisms.

Reprints will not be available from the authors.

The authors declare no conflicts of interest

DISCLOSURES

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Contribution: This author helped design the study, conduct the study, analyze the data, and write the manuscript **Attestation:** Robert J. Brosnan has seen the original study data, reviewed the analysis of the data, approved the final manuscript, and is the author responsible for archiving the study files

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Introduction

Additive anesthetic interactions have been used to support a common mechanism of anesthetic action, although this inference may not apply when receptor occupancy at sites mediating immobility is much less than the mean effective concentration (EC_{50}) for anesthetics at individual sites.¹ In studies *in vivo*, inhaled anesthetic effects are generally additive, even when the combinations of drugs are postulated to exert their effects via different cell mechanisms.^{2,3} In contrast, injectable anesthetics in combination with inhaled anesthetics often produce synergistic effects.^{4,5}

These findings reflect *in vitro* responses of many anesthetic-sensitive ion channels. Mixtures of halothane and isoflurane produce additive responses on gamma-aminobutyric acid type A (GABA_A) and glycine receptors expressed in oocytes.⁶ Despite marked differences in N-methyl-D-aspartate (NMDA) receptor effects at a minimum alveolar concentration, the volatile anesthetics benzene and isoflurane exhibit additive effects on expressed NMDA receptor currents.⁶ Nevertheless, the injectable anesthetic ketamine combined with either isoflurane, sevoflurane, or desflurane synergistically inhibits NMDA receptor currents.⁷

At issue is whether anesthetics acting through different mechanisms must show synergy ⁸, or conversely, whether it is possible to infer a mechanism of anesthetic action based on an additive interaction. To address this question *in vitro*, it is essential to study drugs with known effects on an anesthetic-sensitive ion channel that is likely relevant to anestheticmediated immobility, such as the NMDA receptor.⁹ Carbon dioxide (CO₂) and hydrogen ions (H⁺) both negatively modulate NMDA receptors through extracellular acidification.¹⁰ Isoflurane does not change solution pH; it competitively inhibits NMDA receptor function at the glycine binding site.¹¹ Ketamine antagonizes NMDA receptors by both open channel block and by closed channel block associated with decreased opening frequency.¹² Ketamine must access at least one of these binding sites via the cell membrane,¹² and subunit mutations that diminish NMDA receptor sensitivity to volatile anesthetics do not affect responses to ketamine.¹³ Thus CO₂ and H⁺, isoflurane, and ketamine offer three pharmacologic tools to inhibit NMDA receptors through three different molecular mechanisms. We hypothesized that drug pairs acting via identical mechanisms, such as CO₂ + protons, would inhibit NMDA receptor currents additively, and that drugs pairs acting via different molecular sites, such as CO₂ + ketamine or CO₂ + isoflurane, would inhibit NMDA receptor currents synergistically.

The first study aim was to determine whether anesthetics that inhibit NMDA receptors differently always synergistically inhibit NMDA currents in combination. CO_2 and H⁺ would serve as a positive control for an additive interaction. Ketamine and either CO_2 or H⁺ would serve as a positive control for a synergistic interaction. The inhaled anesthetics CO_2 and isoflurane, having different molecular mechanisms, were hypothesized to inhibit NMDA receptors synergistically. As a second study aim, the effects of CO_2 on the glutamate and glycine binding sites of NMDA receptors would be determined and compared to known isoflurane interactions at these sites.

Methods

Oocyte Preparation and Receptor Expression

An ovary from a tricaine-anesthetized adult female *Xenopus laevis* frog was removed surgically according to a protocol approved by the Animal Care and Use Committee at the University of California, Davis. After defolliculation in a 0.2% Type I collagenase solution (Worthington Biochemical, Lakewood, NJ), oocytes were stored in a filtered modified Barth's solution composed of 88 mM NaCl, 1 mM KCl, 2.4 mM NaHCO₃, 20 mM HEPES,

0.82 mM MgSO₄, 0.33 mM Ca(NO₃)₂, 0.41 mM CaCl₂, 5 mM sodium pyruvate, gentamycin, penicillin, streptomycin, and corrected to pH=7.4. All salts and antibiotics were A.C.S. grade (Fisher Scientific, Pittsburgh, PA).

The human GluN1 NMDA subunit cloned in a pCDNA plasmid and the rat NMDA GluN2A subunit cloned in a pBSII KS+ plasmid were made available by a generous gift from Dr. Adron Harris (University of Texas, Austin). After plasmid linearization, cRNA was synthesized using a T7 transcription kit (mMessage mMachine, Ambion, Austin, TX). Equal proportions of each subunit cRNA were mixed and diluted with DEPC-treated water to 1 mg/mL total nucleic acid. A microinjection pipette (Nanoject II, Drummond Scientific, Broomall, PA) was used to deliver 9 or 18 nL of the transcript mixture or water (controls). Electrophysiology studies on oocytes were conducted 1 or 2 days later.

Voltage Clamp Protocol

Electrophysiology techniques are similar to published protocols.^{10,14–16} Oocytes were studied in a 250µL linear-flow perfusion chamber through which syringe pumps (Pump 33, Harvard Apparatus, Holliston, MA) delivered 1.5 ml/min of barium frog Ringer's solution (BaFR) consisting of 115mM NaCl, 2.5mM KCl, 1.8mM BaCl₂, 10mM HEPES, 0.1 mM EGTA, filtered, and corrected to pH=7.4. Syringes and tubing were made only of glass and PTFE to prevent plasticizer contamination.¹⁷ A –80mV membrane potential was maintained using a standard two-electrode voltage clamp technique (GeneClamp 500B, Axon Instruments, Union City, CA). After a 5 min baseline measurement during perfusion with BaFR, the perfusate was switched to an agonist solution (BaFREG) composed of BaFR plus 0.1 mM glutamate (E) plus 0.01 mM glycine (G) for 30 seconds followed by a 5 min washout with BaFR. This was repeated 3–4 times to verify constancy of the control agonist response (<10% change in peak current). BaFREG itself produced cell currents ≥ 98% of a maximal response.

The perfusate was then switched to the anesthetic test solution in BaFR for a 2 min washin, followed by a 30 sec exposure to the anesthetic test solution in BaFREG. The anesthetic was then washed out with BaFR for 5 min followed by another 30 sec exposure to BaFREG. Data were only used for analysis if NMDA receptor current responses after washout differed by $\leq 10\%$ from responses before washin of the test solution. Data were recorded by commercially available data acquisition software (Chart, version 5, AD Instruments, Colorado Springs, CO).

Anesthetic Concentration-Response Measurement

All test solutions were prepared in 100ml gastight glass syringes using 50ml of either oxygenated BaFR or oxygenated BaFREG in each syringe; drug concentrations were equal in paired syringes. Carbon dioxide test solutions were prepared by using 99.999% pure CO₂ (Matheson Trigas, Newark, CA), and PCO₂ was measured using an automated and calibrated gas analyzer with temperature correction to 23°C (ABL5, Radiometer America, Westlake, OH). The pH test solutions were prepared by adding 1M HCl, and pH was measured using a calibrated meter (Accumet XL20, Fischer Scientific, Hampton, NH). Isoflurane solutions were prepared by adding to the solutions 50ml of a desired isoflurane-oxygen mixture collected downstream from 2 precision out-of-circuit isoflurane vaporizers, shaking vigorously for 30sec, expelling all of the headspace gas, and repeating headspace exchanges 9 additional times. The final isoflurane headspace concentration was measured after a 1.3 min retention time by a calibrated gas chromatograph (Clarus 500, Perkin Elmer, Waltham, MA) with a 0.25ml sample loop, 3m SF-96 packed column with Chromasorb WHP support (Perkin Elmer, Waltham, MA), 35ml/min hydrogen flow, 350 ml/min zero air flow, and 150°C oven temperature. Some isoflurane headspace concentrations were also

measured using an infrared anesthetic gas analyzer (Andros 4800, LumaSense, Santa Clara, CA) that was calibrated using gas chromatography. A 1/4 atmosphere isoflurane concentration was prepared by direct anesthetic injection into fully oxygenated perfusates within gastight syringes. Solutions containing racemic ketamine (Sigma, St. Louis, MO) were prepared by serial dilution.

NMDA responses were measured as a percent of baseline agonist response. Full drug concentration-response curves for NMDA receptor currents were obtained for CO₂, isoflurane, and ketamine. Hill equations were fit to the data using nonlinear regression (v.11, SPSS, Chicago, IL). Since CO₂ effects on NMDA receptor current are mediated by pH ¹⁰, the responses to H⁺ were only measured next to the predicted half-maximal response to confirm the mean inhibitory concentration (IC₅₀) using a simple linear regression model, and maximal inhibition efficacy was assumed.

To test for additivity, a solution containing each study compound was prepared at its respective IC₅₀. Equal volumes of solutions pairs were then combined to yield one-half IC₅₀ concentrations for each compound. The effect of binary drug combinations on NMDA receptor currents in voltage-clamped oocytes was measured, and a 50% current inhibition defined an additive drug interaction. Synergy was defined as an interaction yielding >50% inhibition with a $P \le 0.05$ using a t-test.

Glycine and Glutamate Effect Site Measurement

The nature of CO₂ interactions at the NMDA receptor's glycine and glutamate binding sites was determined by measuring whole cell NMDA receptor current responses in oocytes as a function of glycine concentration or glutamate concentration, respectively. Responses were measured in solutions containing either 0 or 17 mmHg of CO₂ in perfusate solutions, as determined at 37°C using an automated blood gas analyzer. Glycine concentration-responses were compared to a maximal 0.1 mM glycine concentration response. Glutamate concentration responses. Hill equations were fit to the data using nonlinear regression, and the maximum current response (E_{max}), median effective agonist concentration (EC₅₀), Hill coefficient (n_H), and respective bootstrap estimates of standard errors¹⁸ were calculated. Differences between model parameters in the presence and absence of CO₂ were deemed significant when P<0.05 using a t-test.

Competitive inhibition by CO_2 at an agonist site was inferred if CO_2 significantly increased the agonist EC_{50} but did not change E_{max} . Noncompetitive inhibition was inferred if CO_2 significantly decreased the agonist E_{max} but did not change the agonist EC_{50} . If CO_2 significantly decreased E_{max} and increased the agonist EC_{50} , mixed inhibition at the agonist site was inferred.¹⁹

Results

NMDA receptor current was inhibited by CO_2 , H⁺, isoflurane, and ketamine. A sample whole cell current response to anesthetic exposure is shown in Figure 1, and concentration-response curves for each test agent are shown in Figure 2.

NMDA receptor responses to binary drug mixtures, with each applied at one-half of its respective IC_{50} , are shown in Table 1. The combination of CO_2 and H⁺ ions, each with a common mechanism of action on NMDA receptor function, inhibited whole cell current by 51%, consistent with a predicted additive interaction (Figure 3). In contrast, the combination of either CO_2 or H⁺ ions with the injectable anesthetic ketamine, which inhibits NMDA receptor function through different mechanisms of action, inhibited whole cell current by

64% to 67%, consistent with a predicted synergistic interaction. However, combinations of either CO₂ or H⁺ with the inhaled anesthetic isoflurane exhibited additive effects (Figure 3). In a separate experiment with 8 oocytes, application of ¹/₄ EC₅₀ CO₂ plus ³/₄ EC₅₀ isoflurane yielded 49 ± 2 % NMDA receptor inhibition, also indicative of additivity.

NMDA receptor concentration-response curves for the agonists glycine and glutamate, with and without an IC_{50} of CO_2 , are shown in Figure 4, and model parameters are given in Table 2. CO_2 caused a decrease in the maximum glutamate efficacy (E_{max}) without changing the glutamate EC_{50} , which is consistent with noncompetitive inhibition by CO_2 at the NMDA receptor glutamate binding site. CO_2 similarly caused a decrease in the glycine E_{max} , but it also increased the glycine EC_{50} . This is consistent with mixed inhibition by CO_2 at the NMDA receptor glycine binding site. Both glutamate and glycine sigmoid effect models had Hill coefficients >1, indicating cooperative agonist binding. However, only glycine exhibited a change in cooperative NMDA receptor binding with CO_2 acidification.

Discussion

 CO_2 and protons act via an identical mechanism to inhibit NMDA receptor currents in an additive manner.¹⁰ CO₂ and ketamine ¹² act via different mechanisms and combine to synergistically inhibit NMDA receptors. Isoflurane is a competitive antagonist at the glycine binding site of NMDA receptors ¹¹, whereas CO₂ is a noncompetitive antagonist at the glutamate binding site and a mixed antagonist at the glycine binding site of NMDA receptors. These inhaled anesthetics, isoflurane and CO₂, nonetheless inhibit NMDA currents additively. Thus, the study hypothesis is shown to be incorrect. Two anesthetics can additively modulate receptor function despite known differences in their molecular mechanisms of action.

Site-directed mutagenesis studies in NMDA receptor subunits suggest that different amino acid regions are responsible for glycine binding and pH sensitivity. Based on three-dimensional projections of the NMDA receptor sequence onto the lysine-argenine-ornathine-binding protein structure, the glycine binding cleft is located towards the C-terminal half of the extracellular loop formed between membrane segments M3 and M4^{20,21} on the GluN1 subunit and contains amino acids that confer volatile anesthetic sensitivity to NMDA receptors.¹³ This site is close to, but distinct from, the extracellular GluN1 regulatory region encoded by exon 5 (S191 to K211) that confers proton sensitivity to NMDA receptors ²², although mutations within the M3-M4 extracellular loop that are proximal to the putative glycine binding site ^{23,24} or that link it to the M3 segement ²⁴ can modulate pH sensitivity. Amino acids remote from the glycine binding site may also affect the proton IC₅₀, including residues within the M2 pore-forming region of the GluN1 subunit.²⁵ Indeed, certain mutations in the GluN2A subunit, which does not bind glycine at all, can markedly diminish pH sensitivity.²⁴

These molecular actions may explain why, despite their different mechanisms, isoflurane and CO_2 combine additively to inhibit NMDA receptor function. Acidification, such as that caused by CO_2 , decreases the opening frequency of neuronal NMDA channels.²⁶ As a result, there is reduced probability that a channel opens and conducts current at least once before either receptor desensitization or ligand dissociation occurs, and thus acidification decreases E_{max} for both glutamate and glycine (Figure 4 and Table 2). However, from these remote sites, protons likely also induce allosteric changes at the glycine binding site on GluN1 that decrease glycine affinity and increase the glycine EC_{50} (Table 2), just as competitive inhibition from isoflurane increases the glycine EC_{50} for NMDA receptors.¹¹ Perhaps, if negative allosteric modulation is the more important mechanism by which CO_2

or H⁺ inhibits NMDA receptors, then additivity between pH and competitive glycine site inhibitors can become possible.

The concentrations of NMDA receptor antagonist drugs used in this study are not all within clinically relevant ranges for several reasons. First, supra-pharmacologic concentrations of ketamine, isoflurane, and protons were used to obtain complete concentration-response curves in order to more accurately fit a sigmoid curve to the data. In contrast, CO₂ was studied at subphysiologic concentrations. This is because oocyte perfusate solutions lack bicarbonate and have a relatively low buffering capacity. Hence, small increases in CO₂ partial pressure cause greater perfusate acidification than would be expected in either blood or cerebrospinal fluid.¹⁰ Second, if clinically relevant concentrations result in low receptor fractional occupancy, it may be possible to observe additive responses even with two drugs acting via two different targets.¹ By testing drug interactions closer to their respective IC_{50} values, low NMDA receptor occupancy is eliminated as a potential explanation for additivity between drugs with different molecular mechanisms. Third, measuring responses at combinations near the IC_{50} , rather than at either extreme of the sigmoid concentrationresponse curve, provides the greatest scale over which to observe any potential drug-drug interaction; namely antagonism, additivity, or synergy. However, since only 1/2 IC₅₀ drug combinations were studied, it is possible that more than one type of interaction might occur using other antagonist drug ratios.

This study has shown that drugs acting on an anesthetic-sensitive ion channel via different molecular mechanisms can combine to act either synergistically, such as with CO₂ plus ketamine, or additively, such as with CO₂ plus isoflurane. This supports previous work in which anesthetics with suspected dissimilar targets nonetheless additively modulated GABA_A, glycine, and NMDA cell receptor currents.⁶ Consequently, a common molecular mechanism of anesthetic action cannot be reliably inferred from additive *in vitro* drug interactions. *In vivo*, inhaled anesthetics generally combine additively to produce a minimum alveolar concentration,³ and this has been taken as evidence for a common site of anesthetic action even though such a site in animals remains elusive.²⁷ However, if *in vivo* responses reflect *in vitro* drug effects, then it may be incorrect to infer mechanisms of anesthetic actions on the basis of additivity alone.

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Figure 1.

Sample physiograph tracing from a two-electrode voltage clamped oocyte expressing N-methyl-D-aspartate (NMDA) receptors. Currents were stimulated using barium frog Ringer's solution (Ba-FR) containing glutamate (E) and glycine (G) Isoflurane reversibly inhibited NMDA receptor current.

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Figure 2.

Concentration effects of CO₂ (A), pH (B), isoflurane (C) and ketamine (D) on N-methyl-Daspartate (NMDA) receptor currents measured using two-electrode voltage clamping. Horizontal error bars denote standard deviation at the IC₅₀. Lines in panels A, C, and D are described by Hill equations (parameters given in Table 1). NMDA receptor responses to isoflurane partial pressures >190 mmHg are extrapolated (dotted line). Brosnan and Pham



Figure 3.

Effects of binary mixtures of N-methyl-D-aspartate (NMDA) receptor inhibitors, each administered at $\frac{1}{2}$ of their respective IC₅₀, on NMDA whole cell currents (n≥6 oocytes tested for each combination). Error bars denote standard errors. Pairs of inhibitors exhibiting additive effects produced a 50% decrease in NMDA receptor current. Ketamine with either CO₂ or H⁺ caused NMDA receptor inhibition that was significantly more than 50%, consistent with a synergistic effect.

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Figure 4.

Glutamate (panel A) and glycine (panel B) agonist concentration effect on N-methyl-Daspartate (NMDA) receptor currents measured using two-electrode voltage clamping in the presence (+CO₂) or absence ($-CO_2$) of 9 mmHg carbon dioxide at 23 °C. Lines are described by Hill equations (parameters given in Table 2), and horizontal error bars denote standard deviations at the EC₅₀. In the glutamate concentration-response curve (panel A), CO₂ only decreases the maximum control current without affecting the glutamate EC₅₀, consistent with noncompetitive inhibition by CO₂ of NMDA receptors at the glutamate binding site. In the glycine concentration-response curve (panel B), CO₂ decreases the maximum control current and increases the glycine EC₅₀, consistent with mixed inhibition by CO₂ of NMDA receptors at the glycine binding site.

Table 1

Maximum N-methyl-D-aspartate (NMDA) receptor inhibition (I_{max}), median inhibitory concentration (IC_{50}) and Hill coefficient (n_H) estimates for agonist response data at 23 °C described by a Hill equation using nonlinear regression analysis. The IC₅₀ for hydrogren ion concentration (H^+) was determined by simple linear regression for pH values close to the IC₅₀; a Hill coefficient for pH effects was therefore not determined (n.d.).

Antagonist	I_{max} (%)	IC ₅₀	n _H
CO ₂ (mmHg)	100 ± 2	9.5 ± 0.4	1.41 ± 0.07
$H^{+}(nM)$	100	173 ± 8	n.d.
Isoflurane (mmHg)	100 ± 5	50.6 ± 4.7	1.25 ± 0.09
Ketamine (µM)	100 ± 2	1.44 ± 0.06	1.34 ± 0.14

Table 2

Maximum N-methyl-D-aspartate (NMDA) receptor effect (E_{max}), median effective concentration (EC_{50}) and Hill coefficient (n_H) estimates for NMDA agonist concentration-responses in the presence (9 mmHg) and absence (0 mmHg) of CO₂. Statistically significant differences between CO₂ treatment parameters for the same agonist (glutamate or glycine) are shaded and boldfaced. A decrease only in the glutamate concentrationresponse E_{max} value in the presence of CO₂ is consistent with noncompetitive inhibition. A decrease in the glycine concentration-response E_{max} with an increase in EC_{50} in the presence of CO₂ is consistent with mixed inhibition.

Agonist	$CO_2(23^\circ C)$	E_{max} (%)	$EC_{50}\left(\mu M\right)$	n _H
Glutamate	0 mmHg	100 ± 1	4.12 ± 0.32	1.30 ± 0.09
	9 mmHg	51 ± 2	4.56 ± 0.90	1.20 ± 0.20
Glycine	0 mmHg	100 ± 1	0.71 ± 0.03	$\textbf{1.44} \pm \textbf{0.07}$
	9 mmHg	51 ± 2	1.68 ± 0.09	2.90 ± 0.48